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strains of the Neisseria genius

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<151> 1998-10-30

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ttg ata gcg aat gcc tac gat gca tgt gct agg gaa gtg gaa gtt agg Leu Ile Ala Asn Ala Tyr Asp Ala Cys Ala Tar Glu yal Glu Val Arg 144 35

tta ttc gat aaa ccg gag cat aaa atc gtt atc tag gat aat ggt ata Leu Phe Asp Lys Pro Glu His Lys Ile Val Ile Lys Asp Asn Gly Ile 50 192 50

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105

gat tat gca gag att cga aga agc aag ggt att tat caa ccg gag ttt Asp Tyr Ala Glu Ile Arg Arg Ser Lys Gly Ile Tyr Gln Pro Glu Phe 130 135

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	-	_	_		ggc Gly							-				624
					acc Thr											672
					tcg Ser 230									-		720
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		_			att Ile		_		-			_	-	_		816
_					act Thr	_	_	_		_			taa			858
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<213> Neisseria meningitidis

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Leu Phe Asp Lys Pro Glu His Lys Ile Val Ile Lys Asp Asn Gly Ile 50 55 60

Gly Met Ser Phe Asp Glu Ile Asn Asp Phe Tyr Leu Arg Ile Gly Arg 65 70 75 80

Asn Arg Arg Glu Glu Lys Gln Ala Ser Pro Cys Gly Arg Ile Pro Thr 85 90 95

Gly Lys Lys Gly Leu Gly Lys Leu Ala Leu Phe Gly Leu Gly Asn Lys
100 105 110

Ile Glu Ile Ser Thr Ile Gln Gly Asn Glu Arg Val Thr Phe Thr Leu 115 120 125

æis.

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Arg 145	Lys	Glu	Ser	Val	Glu 150	Ser	Asn	Ile	Glu	Ser 155	Gly	Thr	Thr	Ile	Thr 160	
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				acc Thr					Ile						-	96
				tta Leu			_	-						_		144
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Gly Glu Arg Val His Leu Val Gln Leu Gln Lys Thr Gly Lys Asn Ala 50 55 60

Leu Asp Phe Tyr Leu Ser Tyr Tyr Leu Gly Gln Ile Thr Ala Thr Asp 65 70 75 80

Pro Asn Ala Gln Ile Gly Ile Leu Ser Arg Asp Gly Gly Tyr Asp Val 85 90 95

Leu Val Glu His Ile Leu Lys Asn His Gln Ala Lys Gly Ile Val Arg 100 105 110

Leu Ala Asn Ile Asp Glu Val Gln His Gln Lys Ile Ala Thr Glu Pro 115 120 125

Pro Ser Ala Leu Leu Glu Asn Thr Pro Gln Pro Glu Thr Thr Leu Lys 130 135 140

Pro Gln Gln Pro Leu Thr Ser Tyr Phe Gln Ala Ala Leu Thr Ala Leu 145 150 155 160

Arg Arg Pro Asp Ala Phe Arg Pro Cys Arg Leu His Asn Leu Arg Gln 165 170 175

Asn Leu Arg Lys His Ile Leu Ser Asp Leu Phe Lys Glu Lys Thr Asp 180 185 190

Glu Glu Cys Glu Ile Thr Thr Ala Asn Val Ile Asn Lys Leu Lys Ala . 195 200 205

Gln Asn Phe Ile Ser Ile Asp Glu Gln Glu Thr Val Ser Tyr His Leu 210 215 220

Ser Asp Asn Asp Leu Leu Gln Arg Ile Gln Arg His Ile Leu Ser Gln 225 230 235 240

Arg Pro Lys Thr Tyr Ala Asp Phe Gln Ala Val Val Gln Asn Arg Ala
245 250 255

Asp Ala Leu His Leu Thr Val Gly Thr Asn Asp Ile Gln Ser Phe Ala 260 265 270

Arg His Leu Arg Asp Gln Asn Leu Ile Arg Gln Asn Asn Gly Lys Ile 275 280 285

Glu Tyr Ala Pro Phe Thr Glu Pro Lys Pro Gln Pro Thr Pro Lys Gln 290 295 300

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Glu Pro Asp Glu Ile Ile Trp Ly ys Val Pro Lys Lys Thr Ala 320 Ile Ala Ala Leu Ser Leu Lys Asn Arg Pro Asn Lys Thr Lys Thr Leu Arg Asn Thr Ile Gln Ala Leu Thr Lys Ser Asn Ala Gln Glu Thr Asp Lys Leu Gln His Leu Gln Asp Asp Pro Ser Pro Thr Tyr 360 · <210> 5 <211> 1572 <212> DNA <213> Neisseria meningitidis <220> <221> CDS <222> (1)..(1569) <400> 5 atg aaa aaa too ott tto gtt oto ttt otg tat toa too ota ott acc 48 Met Lys Lys Ser Leu Phe Val Leu Phe Leu Tyr Ser Ser Leu Leu Thr gcc agc gaa atc gcc tat cgc ttt gta ttc gga att gaa acc tta ccg 96 Ala Ser Glu Ile Ala Tyr Arg Phe Val Phe Gly Ile Glu Thr Leu Pro gct gca aaa atg gca gaa acg ttt gcg ctg aca ttt atg att gct gcg 144 Ala Ala Lys Met Ala Glu Thr Phe Ala Leu Thr Phe Met Ile Ala Ala 35 ctg tat ctg ttt gcg cgt tat aag gct tcg cgg ctg ctg att gcg gtg 192 Leu Tyr Leu Phe Ala Arg Tyr Lys Ala Ser Arg Leu Leu Ile Ala Val ttt ttc gcg ttc agc att att gcc aac aat gta cat tat gcg gtt tat 240 Phe Phe Ala Phe Ser Ile Ile Ala Asn Asn Val His Tyr Ala Val Tyr caa agt tgg atg acg ggc atc aat tat tgg ctg atg ctg aaa gag att 288 Gln Ser Trp Met Thr Gly Ile Asn Tyr Trp Leu Met Leu Lys Glu Ile acc gaa gtc ggc agt gcg ggc gcg tcg atg ttg gat aag ttg tgg ctg 336 Thr Glu Val Gly Ser Ala Gly Ala Ser Met Leu Asp Lys Leu Trp Leu 100 105 cct gcg ttg tgg ggc gtg ttg gaa gtc atg ttg ttt tgc agc ctt gcc 384 Pro Ala Leu Trp Gly Val Leu Glu Val Met Leu Phe Cys Ser Leu Ala 115 120 125 aag tto cac cgt aag acg cat ttt tot goo gat ata ctg ttt goo tto 432 Lys Phe His Arg Lys Thr His Phe Ser Ala Asp Ile Leu Phe Ala Phe 130 135 cta atg ctg atg att ttc gtg cgt tcg ttc gac acg aaa caa gag cac 480 Leu Met Leu Met Ile Phe Val Arg Ser Phe Asp Thr Lys Gln Glu His 145 150 155

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Gly	Ile	Ser	Pro	Lys 165	_	Thr	Tyr	Ser	Arg 170	Ile	Lys	Ala	Asn	Tyr 75	Phe	
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	_	agg Arg 195			-		_	_		-		-				624
		agt Ser														672
		cat His	_	_	-					_	-		_	_		720
		cgg Arg	_	-		_	-		_	_					-	768
		gca Ala			-		-			_		-				816
		ccg Pro 275		_			_	-			_			-		864
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Pro	Ala	Leu 115	Trp	Gly	Val	Leu	Glu 120	Val	Met	Leu	Phe	Cys 125	Ser	Leu	Ala	
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Gly	Ile	Ser	Pro	Lys 165	Pro	Thr	Tyr	Ser	Arg 170	Ile	Lys	Ala	Asn	Tyr 175	Phe	
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Ser Phe Gly Tyr Phe Val Gly Arg Val Leu Pro Tyr Gln Leu Phe Asp

Leu Ser Arg Ile Pro Phe Lys Gln Pro Ala Pro Ser Ly 200 Gln Gly Ser Val Gln Asn Ile Val Leu Ile Met Gly Glu Ser Glu Ser 215 Ala Ala His Leu Lys Leu Phe Gly Tyr Gly Arg Glu Thr Ser Pro Phe Leu Thr Arg Leu Ser Gln Ala Asp Phe Lys Pro Ile Val Lys Gln Ser Tyr Ser Ala Gly Phe Met Thr Ala Val Ser Leu Pro Ser Phe Phe Asn Ala Ile Pro His Ala Asn Gly Leu Glu Gln Ile Ser Gly Gly Asp Thr Asn Met Phe Arg Leu Ala Lys Glu Gln Gly Tyr Glu Thr Tyr Phe Tyr Ser Ala Gln Ala Glu Asn Glu Met Ala Ile Leu Asn Leu Ile Gly Lys 305 Lys Trp Ile Asp His Leu Ile Gln Pro Thr Gln Leu Gly Tyr Gly Asn 330 Gly Asp Asn Met Pro Asp Glu Lys Leu Leu Pro Leu Phe Asp Lys Ile 345 Asn Leu Gln Gln Gly Arg His Phe Ile Val Leu His Gln Arg Gly Ser His Ala Pro Tyr Ser Ala Leu Leu Gln Pro Gln Asp Lys Val Phe Gly 375 Glu Leu Ile Val Asp Lys Tyr Asp Asn Thr Ile His Lys Thr Asp Gln 395 Met Ile Gln Thr Val Phe Glu Gln Leu Gln Lys Gln Pro Asp Gly Asn 405 Trp Leu Phe Ala Tyr Thr Ser Asp His Gly Gln Tyr Val Arg Gln Asp 425 Ile Tyr Asn Gln Gly Thr Val Gln Pro Asp Ser Tyr Leu Val Pro Leu 435 Val Leu Tyr Ser Ser Asn Lys Ala Val Gln Gln Ala Ala Asn Gln Ala 455 Phe Ala Pro Cys Glu Ile Ala Phe His Gln Gln Leu Ser Thr Phe Leu 470 465 475 Ile His Thr Leu Gly Tyr Asp Met Pro Val Ser Gly Cys Arg Glu Gly 485 490 Ser Val Thr Gly Asn Leu Ile Thr Gly Asp Ala Gly Ser Leu Asn Ile 505 Arg Asp Gly Lys Ala Glu Tyr Val Tyr Pro Gln 515 520

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	c ccc gaa cto e Pro Glu Leo 150	Tyr Gly A	-		-
	a aac tat acc s Asn Tyr Th: 165				
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					tac Tyr											912
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_		_				aac Asn 535		_	-	_	-	_	-			1632
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	gct Ala 770				-	_	_		-	_		_	_	-	_	2352
	cgc Arg															2400
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	gcc Ala					_		-		_	4-			_		2544
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	ctc Leu															2928
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Tyr Thr Arg Leu Gly. Leu Leu Lys Val Asp Gly Thr Al 600 Gly Gly Lys Leu Tyr Met Ser Ala Arg Gly Lys Gly Ala Gly Tyr Leu Asn Arg Thr Gly Gln Arg Val Pro Phe Leu Ser Ala Ala Lys Ile Gly Arg Asp Tyr Ser Phe Phe Thr Asn Ile Glu Thr Asp Gly Gly Leu Leu Ala Ser Leu Asp Ser Val Glu Lys Thr Ala Gly Ser Glu Gly Asp Thr Leu Ser Tyr Tyr Val Arg Arg Gly Asn Ala Ala Arg Thr Ala Ser Ala Ala Ala His Ser Ala Pro Ala Gly Leu Lys His Ala Val Glu Gln Gly 695 Gly Ser Asn Leu Glu Asn Leu Met Val Glu Leu Asp Ala Ser Glu Ser Ser Ala Thr Pro Glu Thr Val Glu Thr Ala Ala Ala Asp Arg Thr Asp Met Pro Gly Ile Arg Pro Tyr Gly Ala Thr Phe Arg Ala Ala Ala Ala Val Gln His Ala Asn Ala Ala Asp Gly Val Arg Ile Phe Asn Ser Leu Ala Ala Thr Val Tyr Ala Asp Ser Thr Ala Ala His Ala Asp Met Gln Gly Arg Arg Leu Lys Ala Val Ser Asp Gly Leu Asp His Asn Ala Thr 785 Gly Leu Arg Val Ile Ala Gln Thr Gln Gln Asp Gly Gly Thr Trp Glu 810 Gln Gly Gly Val Glu Gly Lys Met Arg Gly Ser Thr Gln Thr Val Gly 825 Ile Ala Ala Lys Thr Gly Glu Asn Thr Thr Ala Ala Ala Thr Leu Gly 840 Met Gly His Ser Thr Trp Ser Glu Asn Ser Ala Asn Ala Lys Thr Asp 850 855 Ser Ile Ser Leu Phe Ala Gly Ile Arg His Asp Ala Gly Asp Ile Gly Tyr Leu Lys Gly Leu Phe Ser Tyr Gly Arg Tyr Lys Asn Ser Ile Ser Arg Ser Thr Gly Ala Asp Glu His Ala Glu Gly Ser Val Asn Gly Thr 905 Leu Met Gln Leu Gly Ala Leu Gly Gly Val Asn Val Pro Phe Ala Ala . 915 Thr Gly Asp Leu Thr Val Glu Gly Gly Leu Arg Tyr Asp Leu Leu Lys 930 935 940

taa

Gln Asp Ala Phe Ala. Lys Gly Ser Ala Leu Gly Trp Se 960 Ser Leu Thr Glu Gly Thr Leu Val Gly Leu Ala Gly Leu Lys Leu Ser Gln Pro Leu Ser Asp Lys Ala Val Leu Phe Ala Thr Ala Gly Val Glu 985 Arg Asp Leu Asn Gly Arg Asp Tyr Thr Val Thr Gly Gly Phe Thr Gly Ala Thr Ala Ala Thr Gly Lys Thr Gly Ala Arg Asn Met Pro His Thr 1010 Arg Leu Val Ala Gly Leu Gly Ala Asp Val Glu Phe Gly Asn Gly Trp 1035 Asn Gly Leu Ala Arg Tyr Ser Tyr Ala Gly Ser Lys Gln Tyr Gly Asn 1050 His Ser Gly Arg Val Gly Val Gly Tyr Arg Phe <210> 9 <211> 339 <212> DNA <213> Neisseria meningitidis <220> <221> CDS <222> (1)..(336) <400> 9 atg aac aca ege ate ate gtt teg get geg tte gtt geg ttg gea tta 48 Met Asn Thr Arg Ile Ile Val Ser Ala Ala Phe Val Ala Leu Ala Leu 10 gca ggt tgc ggc tca atc aat aat gta acc gtt tcc gac cag aaa ctt 96 Ala Gly Cys Gly Ser Ile Asn Asn Val Thr Val Ser Asp Gln Lys Leu 20 cag gaa cgt gcc gcg ttt gcc ttg ggc gtc agc caa aat gcc gta aaa 144 Gln Glu Arg Ala Ala Phe Ala Leu Gly Val Ser Gln Asn Ala Val Lys atc agc aac cgc agc aat gaa agc ata cgc atc aac ttt acc gca act 192 Ile Ser Asn Arg Ser Asn Glu Ser Ile Arg Ile Asn Phe Thr Ala Thr 50 240 gtg ggt aag cgc gtg agc caa tgc tat gtt acc agt gta atc agc aca Val Gly Lys Arg Val Ser Gln Cys Tyr Val Thr Ser Val Ile Ser Thr 288 atc ggc gtt acc act tcc gat gca att tgt ttg gga ggc gga acg cac Ile Gly Val Thr Thr Ser Asp Ala Ile Cys Leu Gly Gly Thr His 85 90 336 aaa ggc aaa agt caa tgc aat gct ttg ctt aaa gcg gca ggc cgt tgc Lys Gly Lys Ser Gln Cys Asn Ala Leu Leu Lys Ala Ala Gly Arg Cys 100 105 110

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.

Ala	Val	Ser	Val 100	Gly	Thr	Val	Leu	Val 105	Ser	Val	Val	Ala	Gln 11	Leu	Asp	
	gcg Ala															384
	tac Tyr 130	_	-	-		-				_	_		_	_	_	432
	tcc Ser	_			_				_	_				-		480
	cgg Arg	_	_	_		_		_	-			-	_			528
	ttg Leu	_	_											-	_	576
	ctg Leu															624
	gcg Ala 210															672
-	gaa Glu	_			_		-	-		_					-	720
	ctg Leu															768
_	ggt Gly		_		_	_	-	_	_			-		_		816
	cgc Arg															864
	gcg Ala 290															912
	att Ile															960
	cat His		_	_	-		-	-			_			_		1008
	gtg Val	_			_			_	-	_	_	_	_	-		1056
	ttc Phe		-		_				_			_	_			1104

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ggc act ttt gtt acg ccg ctt tcg ccg gct ttg tgg atg ggc ttg ggt Gly Thr Phe Val Thr Pro Leu Ser Pro Ala Leu Trp Met Gly Leu Gly 385 390 395 400	7
ttg gcg aaa ttg tcg atg ggc aaa cac atc cgt tat tcg ttt ttt tgg Leu Ala Lys Leu Ser Met Gly Lys His Ile Arg Tyr Ser Phe Phe Trp 405 410 415	
gcg tgg ggt ttg tcg ctg gcg ata ttg gcc agt tcg ata gcg gca gga Ala Trp Gly Leu Ser Leu Ala Ile Leu Ala Ser Ser Ile Ala Ala Gly 420 425 430	
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Leu Phe Gly Ala Leu Leu Ala Gly Phe Asp Val Ser Gln Leu Lys Glu 35 40 45	1
Phe Tyr Ser Gly Gly Thr Lys Ser Val Met Gln Ile Val Ile Met Phe 50 55 60	2
Met Phe Ser Ile Leu Phe Phe Gly Ile Met Asn Asp Val Gly Leu Phe 65 70 75 80	
Arg Pro Met Ile Gly Gly Leu Ile Lys Leu Thr Arg Gly Asn Ile Val	
Ala Val Ser Val Gly Thr Val Leu Val Ser Val Val Ala Gln Leu Asp 100 105 110)
Gly Ala Gly Ala Thr Thr Phe Leu Leu Val Val Pro Ala Leu Leu Pro 115 120 125	
Leu Tyr Lys Arg Leu His Met Asn Pro Tyr Leu Leu Phe Leu Leu 130 135 140	1
Thr Ser Ser Ala Gly Leu Ile Asn Leu Leu Pro Trp Gly Gly Pro Thi 145 150 155 160	
Gly Arg Val Ala Ser Val Leu Gly Ala Asp Val Gly Glu Leu Tyr Lys 165 170 175	;
Pro Leu Leu Thr Val Gln Ile Ile Gly Val Val Phe Ile Leu Ala Leu 180 185 190	1
Ser Leu Leu Gly Val Arg Glu Lys Arg Arg Ile Val Arg Glu Leu 195 200 205	1

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Glu 225	Glu	Glu	Gln	Lys	Leu 230	Ala	Arg	Pro	Lys	Leu 235	Phe	Trp	Trp	Asn	Val 240	
Leu	Leu	Phe	Leu	Ala 245	Ala	Met	Ser	Leu	Leu 250	Phe	Ser	Gly	Ile	Phe 255	Pro	
Pro	Gly	Tyr	Val 260	Phe	Met	Leu	Ala	Ala 265	Thr	Ala	Ala	Leu	Leu 270	Leu	Asn	
Tyr	Arg	Ser 275	Pro	Gln	Glu	Gln	Met 280	Glu	Arg	Ile	Tyr	Ala 285	His	Ala	Gly	
Gly	Ala 290	Val	Met	Met	Ala	Ser 295	Ile	Ile	Leu	Ala	Ala 300	Gly	Thr	Phe	Leu	
Gly 305	Ile	Leu	Lys	Gly	Ala 310	Gly	Met	Leu	Asp	Ala 315	Ile	Ser	Lys	Asp	Ile 320	
Val	His	Ile	Leu	Pro 325	Asp	Ala	Leu	Leu	Pro 330	Tyr	Leu	His	Ile	Ala 335	Ile	
Gly	Val	Leu	Gly 340	Ile	Pro	Leu	Glu	Leu 345	Val	Leu	Ser	Thr	Asp 350	Ala	Tyr	
Tyr	Phe	Gly 355	Leu	Phe	Pro	Ile	Val 360	Glu	Gln	Ile	Thr	Ser 365	Gln	Ala	Gly	
Val	Ala 370	Pro	Glu	Ala	Ala	Gly 375	Tyr	Ala	Met	Leu	Ile 380	Gly	Ser	Ile	Val	
Gly 385	Thr	Phe	Val	Thr	Pro 390	Leu	Ser	Pro	Ala	Leu 395	Trp	Met	Gly	Leu	Gly 400	
Leu	Ala	Lys	Leu	Ser 405	Met	Gly	Lys	His	Ile 410	Arg	Tyr	Ser	Phe	Phe 415	Trp	
Ala	Trp	Gly	Leu 420	Ser	Leu	Ala	Ile	Leu 425	Ala	Ser	Ser	Ile	Ala 430	Ala	Gly	
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)> L> CI 2> (3		(1152	2)												
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				caa Gln												96

	gaa Glu	_	-				tat Tyr 40					_	gca Ala	144
	cca Pro 50													192
	aaa Lys	-	-		_	_	_	_	_	_			_	240
	cgt Arg					-				_	_			288
	tta Leu													336
	cgt Arg													384
	ctt Leu 130						-					 _		432
_	gat Asp			-		_			_		-			480
	gcc Ala													528
	gat Asp					_		_	_	_			_	576
	ggt Gly													624
	ctc Leu 210												ccc Pro	672
	ggc Gly													720
	agt Ser													768
	aac Asn													816
	tac Tyr													864
	gct Ala				_	_					-	_		912

290		295		300		
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gaa aag gga ta Glu Lys Gly Ty						
ttg ggg aaa aa Leu Gly Lys Ly 34	ys Val Leu (-		Asn Glu	_	-
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Leu Glu Ala Al 35	la Ser Cys :	Lys Tyr Ile 40	Asn Glu	Ile Tyr 45	Gln Arg	Ala
Asp Pro Thr Al	la Pro Leu :	Phe His Leu 55			Ala Ile	Val
Pro Lys Glu Gl 65	lu Tyr Val (70	Glu Ser Phe	Asp Asp 75	Leu Gly	Lys Thr	Arg 80
Tyr Arg Phe Il	le Lys Ser '	Val Ile Tyr	Glu His 90	Met Lys .	Asn Gly 95	Ala
Ser Leu Val Ty	yr Asn His OO	Ile Asn Asn 105	Glu Pro		Asp His 110	Ile
Ala Arg Gln Va 115	al Ala Arg	Phe Ala Gly 120	Ala His	Thr Ile 125	Val Ser	Gly
Tyr Leu Ala Ph 130	_	Asp Glu Ser 135	_	Asn His	Trp Asp	Thr
Arg Asp Val Ty 145	yr Ala Ile (150	Gln Leu Phe	Gly Lys 155	Lys Arg	Trp Gln	Leu 160
Thr Ala Pro As	sp Phe Pro I 165	Met Pro Leu	Tyr Met 170	Gln Gln	Thr Lys 175	Asp
Thr Asp Ile Se	er Ile Pro (30	Glu His Ile 185	Asp Met		Ile Leu 190	Glu

Ala Gly Asp Val Leu. Ile Pro Arg Gly Trp Trp His Ar Pro Leu Gly Cys Glu Thr Phe His Phe Ala Val Gly Thr Phe Pro Pro Asn Gly Tyr Asn Tyr Leu Glu Trp Leu Met Lys Lys Phe Pro Thr Ile 235 Glu Ser Leu Arg His Ser Phe Ser Asp Trp Glu Gln Asp Arg Thr Arg Ile Asn Asp Thr Ala Ala Gln Ile Ala Ala Met Ile Ala Asp Pro Val Asn Tyr Glu Ala Phe Ser Glu Asp Phe Leu Gly Lys Glu Arg Thr Asp Thr Ala Phe His Leu Glu Gln Phe Ala Asn Pro Asn Ala Thr Pro Leu 295 Ser Asp Asp Val Arg Leu Arg Leu Asn Ala Asn Asn Leu Asp Thr Leu 315 Glu Lys Gly Tyr Leu Ile Gly Asn Gly Met Lys Ile Ser Val Asp Glu 335 Leu Gly Lys Lys Val Leu Glu His Ile Gly Lys Asn Glu Pro Leu Leu 345 Leu Lys Asn Leu Leu Val Asn Phe Asn Gln Gly Lys His Glu Glu Val 355 Arg Lys Leu Ile Tyr Gln Leu Ile Glu Leu Asp Phe Leu Glu Leu Leu 370 375 380 <210> 15 <211> 717 <212> DNA <213> Neisseria meningitidis <220> <221> CDS <222> (1)..(714) <400> 15 atg aat aga ccc aag caa ccc ttc ttc cgt ccc gaa gtc gcc gtt gcc 48 Met Asn Arg Pro Lys Gln Pro Phe Phe Arg Pro Glu Val Ala Val Ala cgc caa acc agc ctg acg ggt aaa gtg att ctg aca cga ccg ttg tca 96 Arg Gln Thr Ser Leu Thr Gly Lys Val Ile Leu Thr Arg Pro Leu Ser ttt tcc cta tgg acg aca ttt gca tcg ata tct gcg tta ttg att atc 144 Phe Ser Leu Trp Thr Thr Phe Ala Ser Ile Ser Ala Leu Leu Ile Ile ctg ttt ttg ata ttt ggt aac tat acg cga aag aca aca gtg gag gga 192 Leu Phe Leu Ile Phe Gly Asn Tyr Thr Arg Lys Thr Thr Val Glu Gly caa att tta cct gca tcg ggc gta atc agg gtg tat gca ccg gat acg 240

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Gln 65	Ile	Leu	Pro	Ala	Ser	Gly	Val	Ile	Arg	Val 75	Tyr	Ala	Pro	Asp	Thr 80	
		att Ile														288
		aag Lys														336
_		cag Gln 115	_	_	_		_		_	-	_	_				384
		cag Gln														432
		ctt Leu														480
		caa Gln														528
		ttg Leu														576
		gaa Glu 195	_	_		-	_	_					_		_	624
		gat Asp	-		-	_	_	-	_		_		-	_		672
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Phe	Ser	Leu 35	Trp	Thr	Thr	Phe	Ala 40	Ser	Ile	Ser	Ala	Leu 45	Leu	Ile	Ile	
Leu	Phe 50	Leu	Ile	Phe	Gly	Asn 55	Tyr	Thr	Arg	Lys	Thr 60	Thr	Val	Glu	Gly	

Gln Ile Leu Pro Ala Ser Gly Val Ile Arg Val Tyr Ala Pro Asp Thr 65 70 75 80

Gly Thr Ile Thr Ala Lys Phe Val Glu Asp Gly Glu Lys Val Lys Ala

Leu Ser Thr Ser Arg Phe Gly Al Gly Asp Lys Leu Phe 105 Ser Val Gln Gln Leu Lys Thr Glu Ala Val Leu Lys Lys Thr Leu 125 Ala Glu Gln Glu Leu Gly Arg Leu Lys Leu Ile His Gly Asn Glu Thr Arg Ser Leu Lys Ala Thr Val Glu Arg Leu Glu Asn Gln Lys Leu His Ile Ser Gln Gln Ile Asp Gly Gln Lys Arg Arg Ile Arg Leu Ala Glu Glu Met Leu Gln Lys Tyr Arg Phe Leu Ser Ala Asn Asp Ala Val Pro 185 Lys Gln Glu Met Met Asn Val Lys Ala Glu Leu Leu Glu Gln Lys Ala Lys Leu Asp Ala Tyr Arg Arg Glu Glu Val Gly Leu Leu Gln Glu Ile 215 Arg Thr Gln Asn Leu Thr Leu Ala Ser Leu Pro Gln Ala Ala 225 230 <210> 17 <211> 690 <212> DNA <213> Neisseria meningitidis <220> <221> CDS <222> (1)..(687) <400> 17 atg atg aat gtc gag gca gag ctt tta gag cag aaa gcc aaa ctt gat 48 Met Met Asn Val Glu Ala Glu Leu Leu Glu Gln Lys Ala Lys Leu Asp 10 gee tae gge ega gaa gaa gee ggg etg ett eag gaa ate ege aeg eag 96 Ala Tyr Gly Arg Glu Glu Ala Gly Leu Leu Gln Glu Ile Arg Thr Gln aat ctg aca ttg gcc agc ctc ccc aaa cgg cat gag aca gaa caa agc 144 Asn Leu Thr Leu Ala Ser Leu Pro Lys Arg His Glu Thr Glu Gln Ser 35 40 cag ctt gaa cgc acc atg gcc gat att tct caa gaa gtt ttg gat ttt 192 Gln Leu Glu Arg Thr Met Ala Asp Ile Ser Gln Glu Val Leu Asp Phe 50 55 gaa atg cgc tct gaa caa atc atc cgt gca gga cgg tcg ggt tat ata 240 Glu Met Arg Ser Glu Gln Ile Ile Arg Ala Gly Arg Ser Gly Tyr. Ile 65 70 gca ata ccg aac gtc gaa gtc gga cag cag gtt gat cct tcc aaa ctg 288 Ala Ile Pro Asn Val Glu Val Gly Gln Gln Val Asp Pro Ser Lys Leu 85 ctc ttg agc att gtt ccc gaa cgt acc gag cta tat gcc cat cta tat

Leu	Leu	Ser	Ile 100	Val	Pro	Glu	Arg	Thr 105	Glu	Leu	Tyr	Ala	His 11	Leu	Tyr	
		agc Ser 115	_	-	_				_	_		-	_	_		384
	_	tat Tyr	_	-									-			432
-	-	gta Val		_	_		_	-	_		-	_	_	_	_	480
	_	ggc Gly	-	_			-	_		_	-		-		-	528
		gtg Val														576
	_	aaa Lys 195	_	_				-	_	_	_	_	_		_	624
		aaa Lys			_		_		-	_		-			-	672
atq	tcq	ggc	aaa	ctq	taa											690
_	-	Gly		_												
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Ser Val	Val	Ser	Val	Ala 150	Lys	Thr	Ala	Leu	Gly 155	Arg	Gln	Glu	Leu	Ser 160	
Gly Leu	ı Gly	Met	Val 165	Ser	Ser	Asp	Leu	Ala 170	Lys	Ser	Asn	Glu	Pro 175	Val	
Tyr Lev	val	Lys 180	Ile	Lys	Pro	Asp	Lys 185	Pro	Thr	Ile	Thr	Ala 190	Tyr	Gly	
Glu Glu	1 Lys 195	Pro	Leu	Gln	Ile	Gly 200	Met	Thr	Leu	Glu	Ala 205	Asp	Ile	Leu	
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aag acq Lys Thr		_										_		_	288
aca gct Thr Ala					_	_						_	_		336
cta caa Leu Glr															384
tcc caa	gct	att	atc	caa	сса	cag	aat	atg	gat	tcg	gga	att	ctg	aaa	432

Ser	Gln 130	Ala	Ile	Ile	Gln	Pro 135	Gln	Asn	Met	Asp	Ser 140	Gly	Ile	Leu	Lys	
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	-	gcg Ala	_		-			_		_						528
		tta Leu									_	-	-		_	576
	_	gaa Glu 195		_	_	_	-	_	_	_			_		_	624
Ile		ccg Pro		-	_	_			_	_		_			J J	672
_	_	aat Asn						_				_	_			720
		acg Thr								-	-		_		-	768
		ttg Leu			-	_	_							_		816
		cac His 275														864
		tcc Ser														912
		ttt Phe														960
		tat Tyr														1008
		gcc Ala														1056
		cga Arg 355														1104
-	_	gaa Glu		_			_	_	_		-			-	-	1152
		cgc Arg														1200

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ata acc gcc gga Ile Thr Ala Gly 435					1344
cag ttt ttc tac Gln Phe Phe Tyr 450					1392
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cag aat act tta Gln Asn Thr Leu 500		_	Asn His Gln I		1536
ggt gcg gac tat Glý Ala Asp Tyr 515					1584
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Ser Met Gln Arg 35	Gln Gln His	Ile Asp Ala 40	Glu Leu Leu 45	Thr Asp Ala	

Asn Val Arg Phe Glu Gln Pro Leu Glu Lys Asn Asn Tyr Val Leu Ser

Thr Arg Val Asn Tyr Ile Ser Leu Glu Asp Glu Thr Pro 70 Lys Thr Ala Arg Lys Phe Ser Phe Leu Pro Ser Val Leu Met Lys Glu 85 Thr Ala Phe Lys Thr Gly Met Cys Leu Gly Ser Asn Asn Leu Ser Arg 105 Leu Gln Lys Ala Ala Gln Gln Ile Leu Ile Val Arg Gly Tyr Leu Thr 125 Ser Gln Ala Ile Ile Gln Pro Gln Asn Met Asp Ser Gly Ile Leu Lys 135 Leu Arg Val Ser Ala Gly Glu Ile Arg Asp Ile Arg Tyr Glu Glu Lys 145 155 Arg Asp Ala Lys Ser Ala Glu Gly Ser Ile Ser Ala Phe Asn Asn Lys 165 170 Leu Pro Leu Tyr Arg Asn Lys Ile Leu Asn Leu Arg Asp Val Glu Gln 180 185 Gly Leu Glu Asn Leu Arg Arg Leu Pro Ser Val Lys Thr Asp Ile Gln Ile Ile Pro Ser Glu Glu Glu Gly Lys Ser Asp Leu Gln Ile Lys Trp 215 Gln Gln Asn Lys Pro Ile Arg Phe Ser Ile Gly Ile Asp Asp Ala Gly 235 230 Gly Lys Thr Thr Gly Lys Tyr Gln Gly Asn Val Ala Leu Ser Ser Asp 245 Asn Pro Leu Gly Leu Ser Asp Ser Phe Tyr Val Ser Tyr Gly Arg Gly 265 Leu Val His Lys Thr Asp Leu Thr Ala Ala Thr Gly Thr Glu Thr Glu Ser Gly Ser Arg Ser Tyr Ser Val His Tyr Ser Val Pro Val Lys Lys 295 Trp Leu Phe Ser Phe Asn His Asn Gly His Arg Tyr His Glu Ala Thr 305 310 315 Glu Gly Tyr Ser Val Asn Tyr Asp Tyr Asn Gly Lys Gln Tyr Gln Ser 325 330 Ser Leu Ala Ala Glu Arg Met Leu Trp Pro Pro Ser Phe Pro Gln Thr 345 Ser Val Arg Met Lys Leu Trp Thr Arg Gln Thr Tyr Lys Tyr Ile Asp 360 Asp Ala Glu Ile Glu Val Gln Arg Arg Ser Ala Gly Trp Glu Ala 375 Glu Leu Arg His Arg Ala Tyr Leu His Arg Trp Gln Leu Asp Gly Lys 390 395 Leu Ser Tyr Lys Arg Gly Thr Gly Met Arg Gln Ser Met Pro Ala Pro

Glu	Glu	Asn	Gly 420	Gly		Thr	Ile	Pro 425	Ala	Thr	Ser	Arg	Ме 430	zys	Ile	
Ile	Thr	Ala 435	Gly	Leu	Asp	Ala	Ala 440	Ala	Pro	Ser	Met	Leu 445	Gly	Lys	Gln	
Gln	Phe 450	Phe	Tyr	Ala	Thr	Ala 455	Ile	Gln	Ala	Gln	Trp 460	Asn	Lys	Thr	Pro	
Leu 465	Val	Ala	Gln	Asp	Lys 470	Leu	Ser	Ile	Gly	Ser 475	Arg	Tyr	Thr	Val	Arg 480	
Gly	Phe	Asp	Gly	Glu 485	Gln	Ser	Leu	Phe	Gly 490	Glu	Arg	Gly	Phe	Tyr 495	Trp	
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Gly	Ala	Asp 515	Tyr	Gly	Arg	Val	Ser 520	Gly	Glu	Ser	Ala	Gln 525	Tyr	Val	Ser	
Gly	Lys 530	Gln	Leu	Met	Gly	Ala 535	Val	Val	Gly	Phe	Arg 540	Gly	Gly	His	Lys	
Val 545	Gly	Gly	Met	Phe	Ala 550	Tyr	Asp	Leu	Phe	Ala 555	Gly	Lys	Pro	Leu	His 560	
Lys	Pro	Lys	Gly	Phe 565	Gln	Thr	Thr	Asn	Thr 570	Val	Tyr	Gly	Phe	Asn 575	Leu	
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Glu Ala Lys Glr · 35	Thr Ala	Leu Ile 40	Ala Thr	Tyr Arg	His Ser 45	Ser Me	t
Val Ala Ala Glu 50	Gln Tyr	Ala Leu 55	Gln Leu	Lys Lys	Ala Gln	Asp Gl	ı
Arg Gln Arg Trp 65	Tyr Asp	Phe Ser	Gln Lys	Gln Gly 75	Arg Lys	Pro Val	
Lys Lys Gln Tyr	Pro Pro 85	Gln Thr	Lys Lys 90	Ala Gly	Tyr Leu	Lys Th	r
Lys Glu Glu Leu 100			Ala Cys 105	Leu Lys	Ala Glu 110	Met Ala	a
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				ggc Gly												192
	-			gct Ala				_							-	240.
				ctg Leu 85												288
		_	-	agg Arg	_		_	_		_	-					336
				tcc Ser			-	_	-	-	_			-		384
				ccg Pro												432
	_	_		cat His		-				_			_			480
				cac His 165												528
		_	_	tcc Ser	_							_		_		576
				cct Pro												624
-	_	_	-	gag Glu		_	_				-		_			672
_		-		ctg Leu			_		-	_		_	-	_		720
		-		ggt Gly 245						-	_					768
_				cta Leu	_		_	-	-		_		-		_	816
gat	gtt	gta	gaa	gtt	gcc	ccc	tct	tac	gac	caa	tcc	gac	att	acc	gct	864

Asp Val Val 275	Glu Val		Ser Tyr 280	Asp	Gln :	Ser As _i 28	4	Thr	Ala	
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Ser His Ala 35	Asp Trp	Val Ile	Thr Gly	Val	Pro :	Tyr Ası 4		Ala	Val	
Ser Gly Arg 50	Ser Gly	Ala Arg 55	Phe Gly	Pro	Glu A	Ala Il 60	e Arg	Arg	Ala	
Ser Val Asn 65	Leu Ala	Trp Glu 70	His Arg	Arg	Phe 175	Pro Tr	Thr	Phe	Asp 80	
Val Arg Glu	Arg Leu 85	Asn Ile	Ile Asp	Cys 90	Gly A	Asp Le	ı Val	Phe 95	Ser.	
Phe Gly Asp	Ser Arg 100	Asp Phe	Val Glu 105	_	Met (Glu Ala	a His 110	Ala	Gly	
Lys Leu Leu 115	Ser Ser	Gly Lys	Arg Cys	Leu	Ser 1	Leu Gl	_	Asp	His	
Phe Ile Thr 130	Leu Pro	Leu Leu 135	Arg Ala	His		Arg Ty:	r Phe	Gly	Lys	
Leu Ala Leu 145	Ile His	Phe Asp 150	Ala His	Thr	Asp 5	Thr Ty	Asp	Asn	Gly 160	
Ser Glu Tyr	Asp His 165	Gly Thr	Met Phe	Tyr 170	Thr A	Ala Pro	Lys	Glu 175	Gly	
Leu Ile Asp	Pro Ser 180	Arg Ser	Val Glr 185		Gly :	Ile Ar	g Thr 190	Glu	His	
Ser Lys Lys 195	Leu Pro	Phe Thr	Val Leu 200	Thr	Ala 1	Pro Gli 20		Asn	Glu	
Asp Ser Val 210	Glu Glu	Thr Val 215	Arg Lys	Ile	_	Glu Th	val	Gly	Asn	
Met Pro Val 225	Tyr Leu	Thr Phe 230	Asp Ile	Asp	Cys 1 235	Leu As _l	Pro	Ser	Phe . 240	
Ala Pro Gly	Thr Gly 245	Thr Pro	Val Cys	Gly 250	Gly 1	Leu Se	s Ser	Asp 255	Arg	

Ala Leu Lys Ile Leu Arg Gly Leu Thr Asp Leu Asp Ile Val Gly Met Asp Val Val Glu Val Ala Pro Ser Tyr Asp Gln Ser Asp Ile Thr Ala 275 Leu Ala Gly Ala Thr Ile Ala Leu Glu Met Leu Tyr Leu Gln Gly Ala 295 300 Lys Lys Asp 305 <210> 25 <211> 426 <212> DNA <213> Neisseria meningitidis <220> <221> CDS <222> (1)..(423) <400> 25 atg gag cag tcg ggc aaa ttc agt tgg tct gcg gca gct ttt tgg gac 48 Met Glu Gln Ser Gly Lys Phe Ser Trp Ser Ala Ala Phe Trp Asp att ccc tac ccc gtc acc agg cgg att gcc tca agt ttg tat tcg acc Ile Pro Tyr Pro Val Thr Arg Arg Ile Ala Ser Ser Leu Tyr Ser Thr 20 gaa tat ttt gtc gta tgc ttt ctg cgt ttg atg cca ctc tct ccg tgt Glu Tyr Phe Val Val Cys Phe Leu Arg Leu Met Pro Leu Ser Pro Cys 35 aat ctg tat ttt gtc acc cat ctg cgt acc aat gaa tcg gaa ata gaa 192 Asn Leu Tyr Phe Val Thr His Leu Arg Thr Asn Glu Ser Glu Ile Glu 50 aga tgg tct gct gtt ccc tgc caa ata gta ttg aac gac ggc aag tcg 240 Arg Trp Ser Ala Val Pro Cys Gln Ile Val Leu Asn Asp Gly Lys Ser 65 gaa ttc ggc gga ttc gca ttt gaa gtg caa ctt tcc cta aca gaa aaa Glu Phe Gly Gly Phe Ala Phe Glu Val Gln Leu Ser Leu Thr Glu Lys ggc cag tat gcg gta gca tac gac ctt tcc tgc aag aaa gat tgc cat 336 Gly Gln Tyr Ala Val Ala Tyr Asp Leu Ser Cys Lys Lys Asp Cys His 100 105 110 gag cta cac gca act gac cca agg cga acg ata cca cat cca ata cct Glu Leu His Ala Thr Asp Pro Arg Arg Thr Ile Pro His Pro Ile Pro 115 120 gtc ccg cca ctg cac cgt cac cga aat cgc caa aca gct taa 426 Val Pro Pro Leu His Arg His Arg Asn Arg Gln Thr Ala 130 135

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Glu	Tyr	Phe 35	Val	Val	Cys	Phe	Leu 40	Arg	Leu	Met	Pro	Leu 45	Ser	Pro	Cys	
Asn	Leu 50	Tyr	Phe	Val	Thr	His 55	Leu	Arg	Thr	Asn	Glu 60	Ser	Glu	Ile	Glu	
Arg 65	Trp	Ser	Ala	Val	Pro 70	Cys	Gln	Ile	Val	Leu 75	Asn	Asp	Gly ·	Lys	Ser 80	
Glu	Phe	Gly	Gly	Phe 85	Ala	Phe	Glu	Val	Gln 90	Leu	Ser	Leu	Thr	Glu 95	Lys	
Gly	Gln	Tyr	Ala 100	Val	Äla	Tyr	Asp	Leu 105	Ser	Cys	Lys	Lys	Asp 110	Cys	His	
Glu	Leu	His 115	Ala	Thr	Asp	Pro	Arg 120	Arg	Thr	Ile	Pro	His 125	Pro	Ile	Pro	
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Ala Lys Thr Arg Asp Pro Met Trp Asp Asn Ala Ala Leu Gln Thr Ser 35 40 45

Glu Gly Val Asn Phe Ile Ala Arg Phe Leu Gly Phe Phe Ser Asp Gly 50 55 60

Glu Tyr Arg Tyr Val Asp Val Leu Gln Pro Asn His Ser Asp Ile Ile 65 70 75 80

Arg Tyr Ser Gly Lys Asp Phe Pro Leu Asn Gln Ile Leu Asn His Ile 85 90 95

His Pro Ala Arg Tyr Ala Val Thr Phe Glu Asn Asn Val Asp Ser Lys
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Leu Arg Arg His 115

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1 5 10 15

tgc ctg aca ctg acg ccc tat ttg caa cat gaa cta ttt tcg gct atg 96 Cys Leu Thr Leu Thr Pro Tyr Leu Gln His Glu Leu Phe Ser Ala Met 20 25 30

aaa tcc tat ttt tcc aaa tat atc cta ccc gtt tca ctt ttt acc ttg 144 Lys Ser Tyr Phe Ser Lys Tyr Ile Leu Pro Val Ser Leu Phe Thr Leu 35 40 45

cca cta tcc ctt tcc cca tcc gtt tcg gct ttt acg ctg cct gaa gca 192
Pro Leu Ser Leu Ser Pro Ser Val Ser Ala Phe Thr Leu Pro Glu Ala
50 60

tgg cgg gcg cag caa cat tcg gct gat ttt caa gcg tcc cat tac 240

Trp 65	Arg	Ala	Ala	Gln	Gln	His	Ser	Ala	Asp	Phe 75	Gln	Ala	Ser	His	Tyr 80	
				gtg Val 85												288
			_	tcc Ser	-		_	_		-	_	_	_		-	336
				cgc Arg												384
			-	gct Ala	-			_			-		_			432
				gca Ala												480
_	_		-	gcc Ala 165	-	_				-			_	_	-	528
	_	_	-	cat His			_			_		_	-	_	-	576
	_		-	gct Ala						_	_			-	_	624
		-	_	aaa Lys				-		-	_	-				672
_	_	-	-	gag Glu					-		_	-		_		720
				agc Ser 245												768
_	_	_		ctg Leu		_	_	_	_		_	_	_	_		816
_	-		-	tta Leu					_			_	_	_		864
-	_		_	agc Ser		_				_	_	_		_	_	912
			_	tct Ser	_		-			_				Tyr		960
			_	aat Asn 325		_								_	_	1008

Val	ggc Gly	gta Val	cag· Gln 340	ttg, Leu	a A	ttg Leu	ccg Pro	ctt Leu 345	tat Tyr	acc Thr	ggc Gly	gga Gly	gaa Glu 350	g	tcg Ser	1056
				_	gcc Ala	-					_			_	_	1104
					cgg Arg											1152
					gcg Ala 390											1200
_	_	_	_	_	ttg Leu		_		_		_					1248
					cgg Arg											1296
_		_	_	_	aaa Lys	_	_		_					-	-	1344
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Met 1	3> Ne 0> 3(Thr	eisse) Leu	Leu	Asn. 5		Met	Ile		10	-	_	_		15		
Met 1 Cys	3> Ne 0> 30 Thr Leu	eisse) Leu Thr	Leu Leu 20	Asn. 5 Thr	Leu	Met Tyr	Ile Leu	Gln 25	10 His	Glu	Leu	Phe	Ser 30	15 Ala	Met	
Met 1 Cys Lys	3> Ne 0> 30 Thr Leu Ser	Leu Thr Tyr 35	Leu Leu 20 Phe	Asn. 5 Thr	Leu	Met Tyr Tyr	Ile Leu Ile 40	Gln 25 Leu	10 His Pro	Glu Val	Leu Ser	Phe Leu 45	Ser 30 Phe	15 Ala Thr	Met Leu	
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115 120 Thr Leu Phe Asp Ala A Lys Phe Ala Gln Tyr Arg Gln Ser 135 Asp Thr Gln Ala Ala Glu Gln Arg Phe Asp Ala Ala Arg Glu Glu Leu Leu Leu Lys Val Ala Glu Ser Tyr Phe Asn Val Leu Leu Ser Arg Asp Thr Val Ala Ala His Ala Ala Glu Lys Glu Ala Tyr Ala Gln Gln Val Arg Gln Ala Gln Ala Leu Phe Asn Lys Gly Ala Ala Thr Ala Leu Asp Ile His Glu Ala Lys Ala Gly Tyr Asp Asn Ala Leu Ala Gln Glu Ile Ala Val Leu Ala Glu Lys Gln Thr Tyr Glu Asn Gln Leu Asn Asp Tyr 235 Thr Asp Leu Asp Ser Lys Gln Ile Glu Ala Ile Asp Thr Ala Asn Leu 245 Leu Ala Arg Tyr Leu Pro Lys Leu Glu Arg Tyr Ser Leu Asp Glu Trp 265 Gln Arg Ile Ala Leu Ser Asn Asn His Glu Tyr Arg Met Gln Gln Leu 275 Ala Leu Gln Ser Ser Gly Gln Ala Leu Arg Ala Ala Gln Asn Ser Arg 295 Tyr Pro Thr Val Ser Ala His Val Gly Tyr Gln Asn Asn Leu Tyr Thr 305 310 315 Ser Ser Ala Gln Asn Asn Asp Tyr His Tyr Arg Gly Lys Gly Met Ser 325 Val Gly Val Gln Leu Asn Leu Pro Leu Tyr Thr Gly Gly Glu Leu Ser 340 Gly Lys Ile His Glu Ala Glu Ala Gln Tyr Gly Ala Ala Glu Ala Gln Leu Thr Ala Thr Glu Arg His Ile Lys Leu Ala Val Arg Gln Ala Tyr 370 375 Thr Glu Ser Gly Ala Ala Arg Tyr Gln Ile Met Ala Gln Glu Arg Val 395 Leu Glu Ser Ser Arg Leu Lys Leu Lys Ser Thr Glu Thr Gly Gln Gln 405 Tyr Gly Ile Arg Asn Arg Leu Glu Val Ile Arg Ala Arg Gln Glu Val 425 Ala Gln Ala Glu Gln Lys Leu Ala Gln Ala Arg Tyr Lys Phe Met Leu 435 Ala Tyr Leu Arg Leu Val Lys Glu Ser Gly Leu Gly Leu Glu Thr Val 450 455 460 Phe Ala Glu

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<211> 231

<212> PRT

<213> Neisseria meningitidis

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Thr Leu Gly Ile Cys Ala Leu Leu Ala Phe Cys Phe Gly Ala Ala Ile

Ala Ser Gly Tyr His Leu Glu Tyr Glu Tyr Gly Tyr Arg Tyr Ser Ala

Val Gly Ala Leu Ala Ser Val Val Phe Leu Leu Leu Leu Ala Arg Gly

Phe Pro Arg Val Ser Ser Val Val Leu Leu Ile Tyr Val Gly Thr Thr 70

Ala Leu Tyr Leu Pro Val Gly Trp Leu Tyr Gly Ala Pro Ser Tyr Gln

Ile Val Gly Ser Ile Leu Glu Ser Asn Pro Ala Glu Ala Arg Glu Phe 100 105

Val Gly Asn Leu Pro Gly Ser Leu Tyr Phe Val Gln Ala Leu Phe Phe 120

Ile Phe Gly Leu Thr Val Trp Arg Tyr Cys Val Ser Gly Gly Val Phe 130 135

Ala Asp Val Lys Asn Tyr Lys Arg Arg Ser Lys Ile Trp Leu Thr Ile 145 150 155

Leu Leu Thr Leu Ile Leu Ser Cys Ala Val Met Asp Lys Ile Ala Ser 165

Asp Lys Asp Leu Arg Glu Pro Asp Ala Gly Leu Leu Leu Asn Ile Phe 180 185

Asp Leu Tyr Tyr Asp Leu Ala Ser Ala Pro Ala Pro Ile Cys Arg Gln 195 200 205

Ala Arg Pro His Phe Gly Ser Ser Lys Ser Val Asn Met Ala Tyr 220

Pro Ser Cys Cys Ala Gln Val 225 230

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<213> Neisseria meningitidis

195

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225

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185

624

672

720

atg cat gtc ggt aaa ggc ggc gag cga acg ttg aca cat ggt gcg tgg

Met His Val Gly Lys Gly Gly Glu Arg Thr Leu Thr His Gly Ala Trp

aag cgt caa agc tac ggc gtg ccg ctg gtt aaa att tcg tcc gat gac

Lys Arg Gln Ser Tyr Gly Val Pro Leu Val Lys Ile Ser Ser Asp Asp

acg cgg cgc gaa atg att aaa gtg agg cgc agc gcg ttt aat ttt tta

Thr Arg Arg Glu Met Ile Lys Val Arg Arg Ser Ala Phe Asn Phe Leu

235

200

215

cgc gga ttc ggc Arg Gly Phe Gly					
gac ggc tat gat Asp Gly Tyr Asp 260	Phe Trp G				
aat aac ctt gcc Asn Asn Leu Ala 275					
tat gcg gga aaa Tyr Ala Gly Lys 290		er Thr Lys	_	Lys Lys	tga
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Asn Leu Ser Leu 35	Pro Gln Th	er Leu Gly 40	Leu Pro Gly	Glu Pro 45	Asn Asn
Asn Ile Val Ser 50		vs Gln Ala 55	Gly Phe Arg		Trp Leu
Ser Asn Gln Gly 65	Met Leu G	y His Phe	Ala Asn Glu 75	Ile Ser	Thr Tyr 80
Ala Leu Arg Ser	Asp Tyr Pi 85	o Trp Phe	Thr Gln Arg	Gly Asp	Tyr Gly 95
Lys Ser Ala Gly		sp Arg Leu 105	Leu Leu Pro	Ala Phe 110	Lys Arg
Val Leu Ile Gly 115	Asn Ala G	y Thr Lys 120	Pro Arg Leu	Ile Val 125	Met His
Leu Met Gly Ser 130	His Ser As		Thr Arg Leu 140		Asp Ala
Arg Arg Phe Gln 145	Tyr Gln Th	nr Glu Lys	Ile Ser Cys 155	Tyr Val	Ser Thr 160
Ile Ala Gln Thr	Asp Lys Pl 165	ne Leu Glu	Asp Thr Val	Lys Ile	Leu Asn 175
Glu Asn Lys Glu 180	_	er Leu Val 185	Tyr Phe Ser	Asp His	Gly Leu
Met His Val Gly 195	Lys Gly G	y Glu Arg 200	Thr Leu Thr	His Gly 205	Ala Trp
Lys Arg Gln Ser 210	Tyr Gly Va		Val Lys Ile 220		Asp Asp
Thr Arg Arg Glu	Met Ile L	s Val Arg	Arg Ser Ala	Phe Asn	Phe Leu

Arg Gly Phe Gly Ser Tip Thr Gly Ile Glu Thr Asp Glu Leu Pro Asp 245 250 255

Asp Gly Tyr Asp Phe Trp Gly Asn Val Pro Asp Val Gln Gly Glu Gly 260 265 270

Asn Asn Leu Ala Phe Ile Asp Gly Leu Pro Asp Asp Pro Ala Pro Trp 275 280 285

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cga ttg gcc acc caa agc ggt ttc aaa gcc atc tat ctg tcc ggc ggc 144
Arg Leu Ala Thr Gln Ser Gly Phe Lys Ala Ile Tyr Leu Ser Gly Gly
35 40 45

ggc gtg gca gcc tgt tct tgc ggt atc cct gat ttg ggc att acc aca 192 Gly Val Ala Ala Cys Ser Cys Gly Ile Pro Asp Leu Gly Ile Thr Thr 50 55

atg gaa gat gtg ctg atc gac gca cga cgc att acg gac aac gtg gat 240 Met Glu Asp Val Leu Ile Asp Ala Arg Arg Ile Thr Asp Asn Val Asp 65 70 75 80

acg cct ctg ctg gtg gac atc gat gtg ggt tgg ggc ggt gca ttc aat 288
Thr Pro Leu Leu Val Asp Ile Asp Val Gly Trp Gly Gly Ala Phe Asn
85 90 95

att gcc cgt acc att cgc aac ttt gaa cgc gcc ggt gtt gca gcg gtt 336
Ile Ala Arg Thr Ile Arg Asn Phe Glu Arg Ala Gly Val Ala Ala Val
100 105 110

cac atc gaa gat cag gta gcg caa aaa cgc tgc ggc cac cgt ccg aac 384 His Ile Glu Asp Gln Val Ala Gln Lys Arg Cys Gly His Arg Pro Asn 115 120 125

aaa gcc att gta tct aaa gat gaa atg gtc gac cgt atc aaa gct gcc 432 Lys Ala Ile Val Ser Lys Asp Glu Met Val Asp Arg Ile Lys Ala Ala 130 135 140

gta gat gcg cgc gtt gat gag aac ttc gtg att atg gcg cgt acc gat 480 Val Asp Ala Arg Val Asp Glu Asn Phe Val Ile Met Ala Arg Thr Asp 145 150 155 160

gcg ctg gcg gta gaa ggt ttg gat gcc gct atc gaa cgc gcc aa gct Ala Leu Ala Val·Glu. Leu Asp Ala Ala Ile Glu Arg Ala h Ala 165	528
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ttg aac atg tac cgc caa ttt gca gat gcg gtg aaa gtg ccc gtg ttg Leu Asn Met Tyr Arg Gln Phe Ala Asp Ala Val Lys Val Pro Val Leu 195 200 205	624
gcg aac att acc gag ttt ggt tcc act ccg ctt tat acc caa agc gag Ala Asn Ile Thr Glu Phe Gly Ser Thr Pro Leu Tyr Thr Gln Ser Glu 210 215 220	672
ctg gct gaa aac ggc gtg tcg ctg gtg ctg tat ccg ctg tca tcg ttc Leu Ala Glu Asn Gly Val Ser Leu Val Leu Tyr Pro Leu Ser Ser Phe 225 230 235 240	· 720
cgt gca gca agc aaa gcc gct ctg aat gtt tac gaa gcg att atg cgc Arg Ala Ala Ser Lys Ala Ala Leu Asn Val Tyr Glu Ala Ile Met Arg 245 250 255	768
gat ggc act tca ggc ggc ggt ggt gga cag tat gca aac ccg tgc cga Asp Gly Thr Ser Gly Gly Gly Gly Gln Tyr Ala Asn Pro Cys Arg 260 265 270	816
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Ala	Leu	Ala	Val	Glu 165	Gly	Leu	Asp	Ala	Ala 170	Ile	Glu	Arg	Ala	Gln 175	Ala	
Cys	Val	Glu	Ala 180	Gly	Ala	Asp	Met	Ile 185	Phe	Pro	Glu	Ala	Met 190	Thr	Asp	
Leu	Asn	Met 195	Tyr	Arg	Gln	Phe	Ala 200	Asp	Ala	Val	Lys	Val 205	Pro	Val	Leu	
Ala	Asn 210	Ile	Thr	Glu	Phe	Gly 215	Ser	Thr	Pro	Leu	Tyr 220	Thr	Gln	Ser	Glu	
Leu 225	Ala	Glu	Asn	Gly	Val 230	Ser	Leu	Val	Leu	Tyr 235	Pro	Leu	Ser	Ser	Phe 240	
Arg	Ala	Ala	Ser	Lys 245	Ala	Ala	Leu	Asn	Val 250	Tyr	Glu	Ala	Ile	Met 255	Arg	
Asp	Gly	Thr	Ser 260	Gly	Gly	Gly	Gly	Gly 265	Gln	Tyr	Ala	Asn	Pro 270	Cys	Arg	
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							_			_		_	act Thr			144
													ggc Gly			192
-				_	_	_		_		_			aac Asn	-	_	240
													tgt Cys			288

gag ctg gcc go Glu Leu Ala Al 115							
atc cgt gtg at Ile Arg Val Il 130							
atg gac gtg at Met Asp Val Me 145						His P	
gaa cgt gaa go Glu Arg Glu Gl							
ctg atc gcc ac Leu Ile Ala Se 18	r Leu Gly						
cac aac ggc aa His Asn Gly Ly 195							
ggt cat ttc ct Gly His Phe Le 210	-	-		_			
atc aaa gcc at Ile Lys Ala Me 225						Glu Pl	
aac gct tct ac Asn Ala Ser Th							
atg tac tcc ag Met Tyr Ser Se 26	r Ile Thr						
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Leu Ser Asp Cy 2	s Leu Ser O	Thr Asn	Pro His 25	Arg Ser	Ser Glu 30	Asn Pa	ro

Lys Pro Ile Lys Thr Glm Arg Arg Asn Thr Met Thr Glu Thr Thr Gln Thr Pro Thr Leu Lys Pro Lys Lys Ser Val Ala Leu Ser Gly Val Ala Ala Gly Asn Thr Ala Leu Cys Thr Val Gly Arg Thr Gly Asn Asp Leu Ser Tyr Arg Gly Tyr Asp Ile Leu Asp Leu Ala Gln Lys Cys Glu Phe Glu Glu Val Ala His Leu Leu Ile His Gly His Leu Pro Asn Lys Phe 105 Glu Leu Ala Ala Tyr Lys Ala Lys Leu Lys Ser Met Arg Gly Leu Pro Ile Arg Val Ile Lys Val Leu Glu Ser Leu Pro Ala His Thr His Pro 135 Met Asp Val Met Arg Thr Gly Val Ser Met Leu Gly Cys Val His Pro 155 Glu Arg Glu Gly His Pro Glu Ser Glu Ala Arg Asp Ile Ala Asp Lys 170 Leu Ile Ala Ser Leu Gly Ser Ile Leu Leu Tyr Trp Tyr Gln Tyr Ser 185 His Asn Gly Lys Arg Ile Glu Val Glu Ser Glu Glu Glu Thr Ile Gly Gly His Phe Leu His Leu Leu His Gly Lys Arg Pro Ser Glu Ser His 210 215 Ile Lys Ala Met His Val Ser Leu Ile Leu Tyr Ala Glu His Glu Phe 230 235 Asn Ala Ser Thr Phe Thr Ala Arg Val Ile Ala Gly Thr Gly Ser Asp 245 Met Tyr Ser Ser Ile Thr Gly Ala Ile Gly Ala Leu Lys Gly Pro Lys 265 His Gly Gly Ala Asn Glu Gly Leu Thr Ile Phe Lys Asn Ala Thr Ala 275 Met Pro Thr Lys Pro Lys Pro Thr Ser Ala Asn Ala Ser Ala Ala Lys 290 295 300 Lys Ser 305 <210> 39 <211> 945 <212> DNA <213> Neisseria meningitidis <220>

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					acc Thr 70											240
	_			_	att Ile	-	_	_	_			-	-		-	288
_					tat Tyr	_	_				_			_		336
					gcg Ala			-	_	-		_	-	_	_	384
_	_		-	_	ata Ile	_	-	_		_		_	_	-		432
					gtc Val 150											480
_					gga Gly	-		_			_		_		-	528
					ggc Gly											576
					ggt Gly			-				_	_			624
					ttt Phe											672
					ttt Phe 230		_			_	-	_			_	720
					ttt Phe						_		_		_	768
					cag Gln											816

260		265	270
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35	4		45
Gln His Ile Ala 50	Phe Leu Lys Th	r Leu Thr Gly Phe 60	Val Phe Ile Ser
Ile Leu Leu Cys	Arg Thr Gly Ph	e Thr Arg Gln Ile	Ala Asp Ile Ser
65	70		80
Arg Lys Lys Glu	Ala Ile Leu Pr	o Leu Leu Lys	Vål Ala Ile Cys
	85	90	95
Ala Phe Phe Gly	Ile Tyr Thr Le	u Phe Phe Phe Glu 105	Thr Thr Ala Tyr 110
Gln Tyr Gly Asn	Ala Ala Asn Va	l Val Val Val Leu	Met Ala Ser Ala
115	12	O	125
Ala Val Ser Ala	Leu Ile Leu As	p Ser Ile Leu Leu	Asp Glu Arg Ile
130	135	140	
Cys Ile Ser Ser	Val Val Gly Va	l Gly Leu Ala Val	Leu Gly Ile Ala
145	150	155	160
Met Ile Ser Trp	Thr Gly Glu Gl	y Ser Leu Gly Leu	Ile Leu Asn Ala
	165	170	175
Ala Leu Ala Gly	Ser Gly Tyr Gl	y Cys Phe Ser Val	Leu Ile Lys Lys
180		185	190
Phe Gly Leu Asn	Gly Gly Ile Ty	r Leu Thr Arg Ile	Leu Met Phe Phe
195	20	O	205
Gly Ser Ile Phe	Leu Phe Ile Pro	o Ser Leu Glu Gly	Ile Glu Asp Ile
210	215	220	
His Trp Gln Trp	Ser Phe Ile Pro	o Pro Leu Leu Ala	Leu Ser Leu Leu
225		235	240

Pro Thr Ile Leu Gly Phe Tyr Cys Thr Thr Lys Ala Leu Asp Tyr Leu Ser Ala Ala Lys Val Gln Val Thr Glu Leu Ala Glu Pro Leu Phe Ala 260 265 Ala Val Leu Ala Trp Leu Phe Leu Asn Glu Ile Pro Glu Gly Arg Phe Phe Val Gly Ala Ile Leu Ile Ile Ala Gly Ile Val Ser Ile Asn Gly 295 Leu Tyr Arg Pro Leu Leu Lys Arg Ile Glu 310 <210> 41 <211> 2610 <212> DNA <213> Neisseria meningitidis <220> <221> CDS <222> (1)..(2607) <400> 41 atg gct gcc aac caa cgt tac cgc aaa ccg ctg ccc ggt acg gat ttg 48 Met Ala Ala Asn Gln Arg Tyr Arg Lys Pro Leu Pro Gly Thr Asp Leu 10 gaa tac tac gac gcg cgt gcg gcg tgt gag gac atc aag ccc ggc tct 96 Glu Tyr Tyr Asp Ala Arg Ala Ala Cys Glu Asp Ile Lys Pro Gly Ser tac gac aag ctg cct tac acg agc cgc att ttg gcg gag aat ttg gtc 144 Tyr Asp Lys Leu Pro Tyr Thr Ser Arg Ile Leu Ala Glu Asn Leu Val aac cgc gcg gac aaa gtc gat ttg ccg acg ctg caa agc tgg ctg ggt 192 Asn Arg Ala Asp Lys Val Asp Leu Pro Thr Leu Gln Ser Trp Leu Gly cag ctg att gag gga aaa cag gaa atc gac ttt cct tgg tat ccg gcg 240 Gln Leu Ile Glu Gly Lys Gln Glu Ile Asp Phe Pro Trp Tyr Pro Ala cgg gtg gtg tgc cac gat att ctg ggg cag acc gcg ttg gtg gat ttg 288 Arg Val Val Cys His Asp Ile Leu Gly Gln Thr Ala Leu Val Asp Leu gca ggt ctg cgc gat gcg att gcc gaa aaa ggc ggc gat cct gcc aaa 336 Ala Gly Leu Arg Asp Ala Ile Ala Glu Lys Gly Gly Asp Pro Ala Lys 100 105 gtg aat ccg gtg gtt gca aaa ccc agc ttc atc gtc gac cac tct ctg 384 Val Asn Pro Val Val Ala Lys Pro Ser Phe Ile Val Asp His Ser Leu 115 120 gee gtt gaa tge gge gge tac gac eee gat gee tte ege aaa aac ege 432 Ala Val Glu Cys Gly Gly Tyr Asp Pro Asp Ala Phe Arg Lys Asn Arg 130 135 caa atc gaa gac aga cgt aac gaa gac cgt ttc cac ttc atc aac tgg 480

Gln Ile Glu Asp Arg Arg Asn Glu Asp Arg Phe His Phe Ile Asn Trp

155

					62-0											528
	-				aat Asn		_		_	_		_	_		-	576
					ttc Phe											624
					gcg Ala											672
_	_	-	-		gtg Val 230	_	_		_			-	_	_	_	720
					gtt Val											768 ·
_		_	_		gtg Val	-	_	_				_	-		_	816
					ttt Phe											864
_				-	cgc Arg						_	_	_			912
					atg Met 310											960
					cgc Arg											1008
					ggc Gly											1056
	Pro				aaa Lys											1104
					ccg Pro											1152
					aaa Lys 390											1200
					atc Ile											1248
tcc	aac	ccg	cgc	aac	gtt	gtt	gcc	gcc	gcg	ctc	ttg	gcg	cgc	aac	gcc	1296

Ser	Asn	Pro	Arg 420	Asn	Val	Val	Ala	Ala 425	Ala	Leu	Leu	Ala	Ar 4	Asn	Ala	
				ctg Leu												1344
				gtg Val												1392
				aaa Lys												1440
				atg Met 485												1488
				gat Asp												1536
				cgt Arg				Tyr								1584
		_	_	gtc Val	_	_		-						-		1632
-		-		gac Asp	-			-	_	_		-	-		_	1680
-		_		tgg Trp 565		-	-	-	-		-	-	-	-	_	1728
_				ccg Pro	_	_		_	-			-	_	_		1776
_				gcg Ala			_		_	_	_		_			1824
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-		_	_	aca Thr		_		_	_	_	_			_		1920
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				ggc Gly												2016
_				tac Tyr	-			-		_		_		-		2064

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ggc gaa acc atg Gly Glu Thr Met					Arg
aaa cag ccg ctc Lys Gln Pro Leu 740					
agc cgc gac tgg Ser Arg Asp Trp 755					
att gtt gcc gaa Ile Val Ala Glu 770		u Arg Ile			
atg ggc gtg ttg Met Gly Val Leu 785		-	_	_	
ctg caa ctg gac Leu Gln Leu Asp					Thr
ccg cgc tgc gac Pro Arg Cys Asp 820	_		-		
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Pro Asp Gly Ala Val Ile Ile Ala Ala Ile Thr Ser Cys Thr Asn Thr

Ser Asn Pro Arg Asn 📶 Val Ala Ala Ala Leu Leu Ala Ar 425 Asn Cys Phe Gly Leu Lys Arg Lys Pro Trp Val Lys Thr Ser Phe Ala Pro Gly Ser Lys Val Ala Glu Ile Tyr Leu Lys Glu Ala Gly Leu Leu 455 Pro Glu Met Glu Lys Leu Gly Phe Gly Ile Val Ala Phe Ala Cys Thr 465 Thr Cys Asn Gly Met Ser Gly Ala Leu Asp Pro Lys Ile Gln Lys Glu Ile Ile Asp Arg Asp Leu Tyr Ala Thr Ala Val Leu Ser Gly Asn Arg 505 Asn Phe Asp Gly Arg Val His Pro Tyr Ala Lys Gln Ala Phe Leu Ala Ser Pro Pro Leu Val Val Ala Tyr Ala Leu Ala Gly Ser Ile Arg Phe 535 Asp Ile Glu Asn Asp Val Leu Gly Val Ala Asp Gly Lys Glu Ile Arg 555 Leu Lys Asp Ile Trp Pro Ala Asp Glu Glu Ile Asp Ala Val Val Ala 565 Glu Tyr Val Lys Pro Gln Gln Phe Arg Asp Val Tyr Val Pro Met Phe 585 Asp Thr Gly Thr Ala Gln Lys Ala Pro Ser Pro Leu Tyr Asp Trp Arg 595 Pro Met Ser Thr Tyr Ile Arg Arg Pro Pro Tyr Trp Glu Gly Ala Leu 615 Ala Gly Glu Arg Thr Leu Arg Gly Met Arg Pro Leu Ala Ile Leu Pro 635 Asp Asn Ile Thr Thr Asp His Leu Ser Pro Ser Asn Ala Ile Leu Ala 650 Val Ser Ala Ala Gly Glu Tyr Leu Ala Lys Met Gly Leu Pro Glu Glu 665 Asp Phe Asn Ser Tyr Ala Thr His Arg Gly Asp His Leu Thr Ala Gln 680 Arg Ala Thr Phe Ala Asn Pro Lys Leu Phe Asn Glu Met Val Lys Asn 690 695 Glu Asp Gly Ser Val Arg Gln Gly Ser Phe Ala Arg Val Glu Pro Glu Gly Glu Thr Met Arg Met Trp Glu Ala Ile Glu Thr Tyr Met Asn Arg 725 Lys Gln Pro Leu Ile Ile Ile Ala Gly Ala Asp Tyr Gly Gln Gly Ser 740 Ser Arg Asp Trp Ala Ala Lys Gly Val Arg Leu Ala Gly Val Glu Ala

Ile Val Ala Glu Gly e Glu Arg Ile His Arg Thr Asn Lea Ile Gly Met Gly Val Leu Pro Leu Gln Phe Lys Pro Asp Thr Asn Arg His Thr Leu Gln Leu Asp Gly Thr Glu Thr Tyr Asp Val Val Gly Glu Arg Thr Pro Arg Cys Asp Leu Thr Leu Val Ile His Arg Lys Asn Gly Glu Thr Val Glu Val Pro Val Thr Cys Arg Leu Asp Thr Ala Glu Glu Val Leu Val Tyr Glu Ala Gly Gly Val Leu Gln Arg Phe Ala Gln Asp Phe Leu Glu Gly Asn Ala Ala 865 <210> 43 <211> 1170 <212> DNA <213> Neisseria meningitidis <220> <221> CDS <222> (1)..(1167) <400> 43 atg ccg caa att aaa att ccc gcc gtt tac tac cgt ggc ggt aca tca 48 Met Pro Gln Ile Lys Ile Pro Ala Val Tyr Tyr Arg Gly Gly Thr Ser aaa ggc gtg ttt ttc aaa cgt tcc gac ctg ccc gag gcg gcg cgg gaa 96 Lys Gly Val Phe Phe Lys Arg Ser Asp Leu Pro Glu Ala Ala Arg Glu 20 25 gcg gga agc gca cgc gac aaa atc ctc ttg cgc gta ctc ggc agc ccg 144 Ala Gly Ser Ala Arg Asp Lys Ile Leu Leu Arg Val Leu Gly Ser Pro gac ccc tac ggc aag cag ata gac ggt ttg ggc aac gcc agt tcg tcc 192 Asp Pro Tyr Gly Lys Gln Ile Asp Gly Leu Gly Asn Ala Ser Ser Ser 55 acc agc aaa gcc gtg att ttg gac aag tcc gaa cgc acc gat cac gat 240 Thr Ser Lys Ala Val Ile Leu Asp Lys Ser Glu Arg Thr Asp His Asp 65 gto gat tac ott tto ggg caa gtt too ato gac aaa oot ttt gto gat 288 Val Asp Tyr Leu Phe Gly Gln Val Ser Ile Asp Lys Pro Phe Val Asp

tgg agt ggc aac tgc ggc aac ctc acc gcc gcc gtg ggc gca ttt gcc

Trp Ser Gly Asn Cys Gly Asn Leu Thr Ala Ala Val Gly Ala Phe Ala

atc gag caa ggc ttg gtc gat aaa tcc aaa atc cct tca gac ggc ccg

Ile Glu Gln Gly Leu Val Asp Lys Ser Lys Ile Pro Ser Asp Gly Pro

120

105

100

115

33.6

384

760

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				acg Thr 165											528
				ggc Gly											576
				gtg Val											624
				ccg Pro		_		_					-		672
_				ttg Leu		_	_				-	_	_	 _	720
				aaa Lys 245											768
	-	-	_	tcc Ser	_	_	_	_	_			_	_	_	816
_		-		ccc Pro	-	_	-			-		_			864
				gac Asp											912
				gcg Ala											960
				ccc Pro 325											1008
				gtg Val											1056
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Ala Gly Ser Ala Arg Asp Lys Ile Leu Leu Arg Val Leu Gly Ser Pro 35 40 45

Asp Pro Tyr Gly Lys Gln Ile Asp Gly Leu Gly Asn Ala Ser Ser Ser 50 55 60

Thr Ser Lys Ala Val Ile Leu Asp Lys Ser Glu Arg Thr Asp His Asp 65 70 75 80

Val Asp Tyr Leu Phe Gly Gln Val Ser Ile Asp Lys Pro Phe Val Asp 85 90 95

Trp Ser Gly Asn Cys Gly Asn Leu Thr Ala Ala Val Gly Ala Phe Ala 100 105 110

Ile Glu Gln Gly Leu Val Asp Lys Ser Lys Ile Pro Ser Asp Gly Pro 115 120 125

Cys Thr Val Lys Ile Trp Gln Lys Asn Ile Gly Lys Thr Ile Ile Ala 130 135 140

His Val Pro Met Gln Asn Gly Ala Val Leu Glu Thr Gly Asp Phe Glu
145 150 155 160

Leu Asp Gly Val Thr Phe Pro Ala Ala Glu Val Gln Ile Glu Phe Leu 165 170 175

Asp Pro Ala Asp Gly Glu Gly Ser Met Phe Pro Thr Gly Asn Leu Val

Asp Glu Ile Asp Val Pro Asn Ile Gly Arg Leu Lys Ala Thr Leu Ile 195 200 205

Asn Ala Gly Ile Pro Thr Val Phe Leu Asn Ala Ala Asp Leu Gly Tyr 210 215 220

Thr Gly Lys Glu Leu Gln Asp Asp Ile Asn Asn Asp Ala Ala Leu 225 230 235 240

Glu Lys Phe Glu Lys Ile Arg Ala Tyr Gly Ala Leu Lys Met Gly Leu 245 250 255

Ile Ser Asp Val Ser Glu Ala Ala Ala Arg Ala His Thr Pro Lys Val 260 265 270

Ala Phe Val Ala Pro Ala Ala Asp Tyr Thr Ala Ser Ser Gly Lys Thr 275 280 285

Val Asn Ala Ala Asp Ile Asp Leu Leu Val Arg Ala Leu Ser Met Gly 290 295 300

اِ

Lys Leu His His Ala N Met Gly Thr Ala Ser Val Ala Ile 320 Ala Ala Ala Val Pro Gly Thr Leu Val Asn Leu Ala Ala Gly Ala Gly Thr Arg Lys Glu Val Arg Phe Gly His Pro Ser Gly Thr Leu Arg Val 345 Gly Ala Ala Ala Glu Cys Gln Asp Gly Gln Trp Thr Ala Thr Lys Ala Val Met Ser Arg Ser Ala Arg Val Met Met Glu Gly Trp Val Arg Val 375 Pro Glu Asp Cys Phe <210> 45 <211> 954 <212> DNA <213> Neisseria meningitidis <220> <221> CDS <222> (1)..(951) <400> 45 atg cgc acg ccg ttt tgt tgg gca tac gcc aat gcc gcc cga ata tcg 48 Met Arg Thr Pro Phe Cys Trp Ala Tyr Ala Asn Ala Arg Ile Ser gca atg ctg ccg gcg tgt tgg gcg cag gcg atg ttg gcc gaa gta atc 96 Ala Met Leu Pro Ala Cys Trp Ala Gln Ala Met Leu Ala Glu Val Ile age tge aac aag get teg teg etg eeg eag eet teg geg aga teg geg 144 Ser Cys Asn Lys Ala Ser Ser Leu Pro Gln Pro Ser Ala Arg Ser Ala ttt aaa tca acc tgc ttc atg ggt gat tct ccg tat ttg gtt cag ata 192 Phe Lys Ser Thr Cys Phe Met Gly Asp Ser Pro Tyr Leu Val Gln Ile gac ttg gtt ttt gcg ccg cag ggc ggt ggc ttc ttt caa gcc gat tat 240 Asp Leu Val Phe Ala Pro Gln Gly Gly Gly Phe Phe Gln Ala Asp Tyr ttt gaa ttt gac ttt gct gcc gaa gcg cac ctg tgc cag cct gcc caa 288 Phe Glu Phe Asp Phe Ala Ala Glu Ala His Leu Cys Gln Pro Ala Gln 85 atc ggc ggc ggc aac ggt agc gat ttt cgg ata acc gcc ggt ggt ttg 336 Ile Gly Gly Asn Gly Ser Asp Phe Arg Ile Thr Ala Gly Gly Leu 100 105 110 ege ate gge cag cag gat aat egg ttt gee gee ggg egg cae etg cae 384 Arg Ile Gly Gln Gln Asp Asn Arg Phe Ala Ala Gly Arg His Leu His 115 120 ggt tcc tgc ctg aac agc gtg gga cag cat ttc caa agg ttg cga cag 432 Gly Ser Cys Leu Asn Ser Val Gly Gln His Phe Gln Arg Leu Arg Gln 130 135 140

ggt Gly 145	cag Gln	cgg Arg	ctg Leu	tcc Ser	150	gaa Glu	gcg Ala	gta Val	gcc Ala	cat His 155	gcg Ala	gtt Val	g A.	tc	gct Ala 160	480
		cgt Arg														528
-	_	ggc			_	_			_		_					576
		cgg Arg 195			_	_					_	_			_	624
		gag Glu														672
		ggc Gly														720
		tcc Ser														768
		cat His				-							_			816
_		gac Asp 275			_	-				_						864
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Ser Cys Asn Lys Ala Ser Ser Leu Pro Gln Pro Ser Ala Arg Ser Ala

Phe Lys Ser Thr Cys Phe Met Gly Asp Ser Pro Tyr Leu Val Gln Ile

Asp Leu Val Phe Ala Pro Gln Gly Gly Phe Phe Gln Ala Asp Tyr

	Phe	Glu	Phe	Asp	.Phe 85		Ala	Glu	Ala	His 90	Leu	Cys	Gln	P	1a 95	Gln	
	Ile	Gly	Gly	Gly 100	Asn	Gly	Ser	Asp	Phe 105	Arg	Ile	Thr	Ala	Gly 110	Gly	Leu	
	Arg	Ile	Gly 115	Gln	Gln	Asp	Asn	Arg 120	Phe	Ala	Ala	Gly	Arg 125	His	Leu	His	
	Gly	Ser 130	Cys	Leu	Asn	Ser	Val 135	Gly	Gln	His	Phe	Gln 140	Arg	Leu	Arg	Gln	
	Gly 145	Gln	Arg.	Leu	Ser	Val 150	Glu	Ala	Val	Ala	His 155	Ala	Val	Ala	Ile	Ala 160	
	Leu	Gln	Arg	Pro	Arg 165	Phe	Pro	Phe	Gln	Ile 170	Gln	Thr	Pro	Phe	Phe 175	Thr	
	Glu	Ser	Gly	Ile 180	Phe	Arg	Arg	Arg	Asn 185	Lys	Val	Asp	Gly	Ile 190	Gly	Lys	
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	Asp	Gly 210	Glu	Ile	Ile	Ala	Phe 215	Leu	Gln	His	Ser	Ala 220	Leu	Met	Ala	Ala	
	Glu 225	Thr	Gly	Phe	Gln	Val 230	Gly	Ala	Ser	Arg	Thr 235	His	His	Phe	Arg	His 240	
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•	Arg	Ile	Asp 275	Arg	Phe	Ala	Val	Gln 280	Phe	Arg	Leu	Ile	His 285	Gly	Thr	Gly	
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_			_		gtg Val			_		-		-	_		_	192
					caa Gln 70											240
					gtg Val											288
					gaa Glu											336
					cat His											384
					ttc Phe											432
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Leu	Asn	Ala	Pro 20	Ser	Glu	Leu	Gly	Lys 25	Gln	Gln	Lys	Leu	Trp 30	Ala	Phe	
Ala	Ala	Ala 35	Leu	Gly	Gln	His	Asp 40	Arg	Ile	Glu	Glu	Val 45	Val	Val	Gly	
	_	_	_							_		_	_			

Met Asn Asn Leu Thr Val Phe Thr Arg Phe Asp Thr Asp Leu Ala Thr 50 55 60

1.2

112 -										75	Thr				Asp 80	
нıs	Gln	Gly	Lys	Leu 85	Val	Glu	Ile	Pro	Val 90	Cys	Tyr	Gly	Gly	Glu 95	Tyr	
Gly	Pro	Asp	Leu 100	Ala	Glu	Val	Ala	Ala 105	Phe	His	Gln	Thr	Val 110	Ile	Ser	
Glu	Ile	Val 115	Arg	Arg	His	Thr	Ala 120	Gln	Thr	Tyr	Thr	Val 125	Phe	Met	Met	
Gly	Phe 130	Gln	Pro	Gly	Phe	Pro 135	Tyr	Leu	Gly	Gly	Leu 140	Pro	Glu	Ala	Leu	
His 145	Thr	Pro	Arg	Arg	Ala 150	Val	Pro	Arg	Thr	Ser 155	Val	Pro	Ala	Gly	Ser 160	
Val	Gly	Ile	Gly	Gly 165	Ser	Gln	Thr	Gly	Val 170	Tyr	Pro	Phe	Ala	Ser 175	Pro	
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Asp	Leu	Asn 195	Pro	Pro	Thr	Leu	Leu 200	Ala	Ala	Gly	Asp	Gln 205	Val	Arg	Phe	
Val	Ala 210	Glu	Arg	Ile	Glu	Pro 215										
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Met 1 gga	Ile	cac His		Ser 5 cac	Ala cgg	Val cgt	Gln	Ala ggc	Pro 10 atc	Ala ggt	His cat	Ile gcc	Gln ggt	Asp 15 gcg	Thr atg	48 <u>.</u> 96
Met 1 gga Gly	cgc Arg	cac His tac Tyr	Val gga Gly	Ser 5 cac His	Ala cgg Arg	Val cgt Arg	Gln tac Tyr ggt	Ala ggc Gly 25 aat	Pro 10 atc Ile att	Ala ggt Gly tta	His cat His	Ile gcc Ala ggc	Gln ggt Gly 30 aac	Asp 15 gcg Ala gac	Thr atg Met gaa	
Met 1 gga Gly gac Asp	cgc Arg acg Thr	cac His tac Tyr gtt Val 35	yal gga Gly 20 gct	Ser 5 cac His ttg Leu atc	Ala cgg Arg gcg Ala	cgt Arg gcg Ala	tac Tyr ggt Gly 40	Ala ggc Gly 25 aat Asn	Pro 10 atc Ile att Ile	Ala ggt Gly tta Leu	His cat His ttg	gcc Ala ggc Gly 45	ggt Gly 30 aac Asn	Asp 15 gcg Ala gac Asp	Thr atg Met gaa Glu ttt	96
Met 1 gga Gly gac Asp ggc Gly	cgc Arg acg Thr acg Thr 50	cac His tac Tyr gtt Val 35 gcc Ala	yal gga Gly 20 gct Ala	Ser 5 cac His ttg Leu atc Ile ccg	Ala cgg Arg gcg Ala gaa Glu ttt	cgt Arg gcg Ala atc Ile 55	tac Tyr ggt Gly 40 gct Ala	Ala ggc Gly 25 aat Asn ttg Leu acc	Pro 10 atc Ile att Ile ggc Gly	Ala ggt Gly tta Leu ggg Gly	His cat His ttg. Leu ata Ile 60	gcc Ala ggc Gly 45 atg Met	ggt Gly 30 aac Asn ctg Leu	Asp 15 gcg Ala gac Asp gtg Val	Thr atg Met gaa Glu ttt Phe gaa	96 144

Gly ggg	cag Gln	acc Thr	ttg Leu 100	aaa Lys		gtg Val	cgt Arg	gcc Ala 105	gtg Val	cag Gln	ggc Gly	atg Met	ta Tyr 110	c	tat Tyr	336
					gga Gly											384
					gcc Ala											432
					ctc Leu 150											480
					ccg Pro											528
					gcc Ala											576
					acg Thr											624
_		-		_	ccg Pro	_		_	-			_	_	-	_	672
					gca Ala 230											720
ccg •Pro					gcc Ala											768
	-		-	-	gcc Ala	_	Āsp	_	Gly		_	-	_		-	816
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Met Ile His Val Ser Ala Val Gln Ala Pro Ala His Ile Gln Asp Thr 1 5 10 15

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Gly Arg Tyr Gly His Arg Tyr Gly Ile Gly His Ala G Asp Thr Val Ala Leu Ala Ala Gly Asn Ile Leu Leu Gly Asn Asp Glu Gly Thr Ala Ala Ile Glu Ile Ala Leu Gly Gly Ile Met Leu Val Phe Glu Arg Asp Thr Pro Phe Cys Leu Thr Gly Ala Val Tyr Gln Ala Glu Leu Asp Gly Glu Pro Val Tyr Ser Tyr Trp Arg Tyr Thr Ala Arg Lys Gly Gln Thr Leu Lys Met Val Arg Ala Val Gln Gly Met Tyr Gly Tyr 105 Val Cys Val Ala Gly Gly Phe Asp Val Pro Glu Val Met Gly Ser Arg 115 Ser Thr Asp Leu Lys Ala Gly Phe Gly Gly His Gln Gly Arg Met Leu Gln Lys Gly Asp Tyr Leu Pro Ile Gly Lys Gly Ala Gln Glu Leu Ser 150 145 155 Lys Val Gly Ile Ala Pro Ile Pro Phe Thr Asp Thr Ile His Leu Val 165 Pro Ser Ser Glu Tyr Ala Ala Phe Ser Glu Lys Gly Arg Leu Asn Leu 185 Glu Arg Glu Thr Trp Thr Leu Gln Ser Asp Ser Asn Arg Met Gly Tyr 200 Arg Phe Asp Gly Gln Pro Leu Thr Leu Ser Gln Pro Leu Glu Met Leu 215 Ser His Ala Val Gln Ala Gly Thr Val Gln Val Pro Pro Gly Gly Lys 230 235 Pro Ile Ile Leu Leu Ala Asp Ala Gln Thr Thr Gly Gly Tyr Pro Lys 245 Ile Ala Thr Val Ala Ala Ala Asp Leu Gly Arg Leu Ala Gln Val Arg 265 ·Phe Gly Ser Lys Val Lys Phe Lys Ile Ile Gly Leu Lys Glu Ala Thr 275 280 Ala Leu Arg Arg Lys Asn Gln Val Tyr Leu Asn Gln Ile Arg Arg Ile 290 295 Thr His Glu Ala Gly

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			-			ctt Leu		_		_	-	_				-	336
	-	_				tta Leu	_	-					_	_			384
J		_		_	_	acc Thr	_									-	432
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						ctg Leu		_	_				_		_	-	576
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						ttg Leu 230											720

4.3

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					ggt Gly											912
					ggt Gly 310											960
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		-	•	-	tcc Ser		-			_		_	-		_	1056
					ctt Leu											1104
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				_	gcg Ala	_			_	_		_	_	_	_	1344
				-	tac Tyr	_		-				-	_			1392
-	_				gaa Glu 470	_	_		_				_			1440
		-	_		ttc Phe	_	-		_	_			_			1488
_	_	_		_	att Ile	_	_	_	_	_	_		-		_	1536

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ccg ctt ttg cag Pro Leu Leu Gln 530	-	-			1632
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aat acg act tgg Asn Thr Thr Trp		=	_		1728
gaa ttt tta aaa Glu Phe Leu Lys 580		-			1776
atg ttt aat acc Met Phe Asn Thr 595	•		-	- -	1824
agt att ccc tat Ser Ile Pro Tyr 610					1872
acc ccg gta gtt Thr Pro Val Val 625					1920
cgg ttg att tgg Arg Leu Ile Trp					1968
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Leu Leu Phe Ser 35	His Ile Gly	Ile Asp Leu 40	Ser Ser Ile 45	Thr Val Ile	

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Lys	His	Gly 435	Ile	Val	Ala	Val	Tyr 440	His	Arg	Asp	Gly	Asp 445	Lys	Val	Val
Tyr	Gly 450	Ala	Asn	Val	Tyr	Asp 455	Gly	Ala	Tyr	Asn	Thr 460	Asp	Val	Phe	Asn
Ser 465	Val	Asn	Gly	Ile	Glu 470	Arg	Ala	Tyr	Leu	Leu 475	Pro	Ser	Leu	Lys	Ser 480
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Arg	Val	Leu	Ser 500	Ala	Ile	Pro	Glu	Met 505	Gln	Ser	Met	Ile	Val 510	Ala	Glu
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Glu	Phe	Leu	Lys 580	Gln	Val	Gln	Ser	His 585	Leu	Thr	Pro	Asp	Gly 590	Ile	Val
		595					Pro 600					605			
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Thr 625	Pro	Val	Val	Phe	Pro 630	Asn	Lys	Glu	Leu	Leu 635	Lys	Gln	Arg	Leu	Ser 640
Arg	Leu	Ile	Trp	Pro 645	Glu	Ser	Gly	Arg	His 650	Val	Phe	Asp	Ser	Ser 655	Thr
Val	Asp	Ala	Ala 660	Ala	Gln	Lys	Val	Val 665	Ser	Arg	Met	Leu	Ile 670	Gln	Met
Thr	Glu	Pro 675	Ser	Ala	Gly	Ala	Glu 680	Val	Ile	Thr	Asp	Asp 685	Asn	Met	Ile
Val	Glu 690	Tyr	Lys	Tyr	Gly	Arg 695	Gly	Ile							

<210> 53 <211> 1040 <212> PRT <213> Neisseria meningitidis

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705

Ser Val Arg Phe Thr Arg Thr Asn Pro Ile Gln Ile Ala Gly The Ser Phe Ser Ala Pro Ile Val Thr Gly Thr Ala Ala Leu Leu Gln Lys Tyr Pro Trp Met Ser Asn Asp Asn Leu Arg Thr Thr Leu Leu Thr Thr 410 Ala Gln Asp Ile Gly Ala Val Gly Val Asp Ser Lys Phe Gly Trp Gly Leu Leu Asp Ala Gly Lys Ala Met Asn Gly Pro Ala Ser Phe Pro Phe Gly Asp Phe Thr Ala Asp Thr Lys Gly Thr Ser Asp Ile Ala Tyr Ser Phe Arg Asn Asp Ile Ser Gly Thr Gly Gly Leu Ile Lys Lys Gly Gly Ser Gln Leu Gln Leu His Gly Asn Asn Thr Tyr Thr Gly Lys Thr Ile Ile Glu Gly Gly Ser Leu Val Leu Tyr Gly Asn Asn Lys Ser Asp Met Arg Val Glu Thr Lys Gly Ala Leu Ile Tyr Asn Gly Ala Ala Ser Gly 515 Gly Ser Leu Asn Ser Asp Gly Ile Val Tyr Leu Ala Asp Thr Asp Arg Ser Gly Ala Asn Glu Thr Val His Ile Lys Gly Asp Leu Gln Leu Gly Gly Glu Gly Thr Leu Tyr Thr Arg Leu Gly Lys Leu Leu Lys Val Asp Gly Thr Ala Met Thr Gly Gly Lys Leu Tyr Met Ser Ala Arg Gly Lys Gly Ala Gly Tyr Leu Asn Arg Thr Gly Gln Arg Val Pro Phe Leu Ser 600 Ala Ala Lys Ile Gly Arg Asp Tyr Ser Phe Phe Thr Asn Ile Glu Thr Asp Gly Gly Leu Leu Ala Ser Leu Asp Ser Val Glu Lys Thr Ala Gly 630 Ser Glu Gly Asp Thr Leu Ser Tyr Tyr Val Arg Arg Gly Asn Ala Ala Arg Thr Ala Ser Ala Ala Ala His Ser Ala Pro Ala Gly Leu Lys His 665 Ala Val Glu Gln Gly Gly Ser Asn Leu Glu Asn Leu Met Val Glu Leu 680 Asp Ala Ser Glu Ser Ser Ala Thr Pro Glu Thr Val Glu Thr Ala Ala 695 700 Ala Asp Arg Thr Asp Met Pro Gly Ile Arg Pro Tyr Gly Ala Thr Phe

710

Arg Ala Ala Ala Val Gln His Ala Asn Ala Ala Asp Gly Val Arg Ile Phe Asn Ser Leu Ala Ala Thr Val Tyr Ala Asp Ser Thr Ala Ala His Ala Asp Met Gln Gly Arg Arg Leu Lys Ala Val Ser Asp Gly Leu Asp His Asn Ala Thr Gly Leu Arg Val Ile Ala Gln Thr Gln Gln Asp Gly Gly Thr Trp Glu Gln Gly Gly Val Glu Gly Lys Met Arg Gly Ser Thr Gln Thr Val Gly Ile Ala Ala Lys Thr Gly Glu Asn Thr Thr Ala Ala Ala Thr Leu Gly Met Gly His Ser Thr Trp Ser Glu Asn Ser Ala Asn Ala Lys Thr Asp Ser Ile Ser Leu Phe Ala Gly Ile Arg His Asp Ala Gly Asp Ile Gly Tyr Leu Lys Gly Leu Phe Ser Tyr Gly Arg Tyr 855 Lys Asn Ser Ile Ser Arg Ser Thr Gly Ala Asp Glu His Ala Glu Gly 870 Ser Val Asn Gly Thr Leu Met Gln Leu Gly Ala Leu Gly Gly Val Asn 885 890 Val Pro Phe Ala Ala Thr Gly Asp Leu Thr Val Glu Gly Gly Leu Arg Tyr Asp Leu Leu Lys Gln Asp Ala Phe Ala Glu Lys Gly Ser Ala Leu 920 Gly Trp Ser Gly Asn Ser Leu Thr Glu Gly Thr Leu Val Gly Leu Ala Gly Leu Lys Leu Ser Gln Pro Leu Ser Asp Lys Ala Val Leu Phe Ala 950 955 Thr Ala Gly Val Glu Arg Asp Leu Asn Gly Arg Asp Tyr Thr Val Thr Gly Gly Phe Thr Gly Ala Thr Ala Ala Thr Gly Lys Thr Gly Ala Arg 985 Asn Met Pro His Thr Arg Leu Val Ala Gly Leu Gly Ala Asp Val Glu Phe Gly Asn Gly Trp Asn Gly Leu Ala Arg Tyr Ser Tyr Ala Gly Ser 1015 Lys Gln Tyr Gly Asn His Ser Gly Arg Val Gly Val Gly Tyr Arg Phe

1030

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	_	ggt Gly	_	_	_		_			-				_		96
		gcg Ala 35														144
		gat Asp		-					-			-				192
	_	agc Ser		_	_			_			_	-				240
		-agg. Arg														288
		aaa Lys														336
	_	atc Ile 115							_		-					384
_		gca Ala					_		-				_			432
_		gag Glu		-					-							480
		gag Glu	_	_						_						528
		ctt Leu		_			-		_	-	_	-				576
	-	agc Ser 195	-			_	_		_			-				624
		aat Asn		_					_		_		_	_		672

gca acc aat att tca tcg tta tct tca aaa ttc gaa cag tat gaa tac 720

Ala 225	Thr	Asn	Ile	Ser	Ser 230	Leu	Ser	Ser	Lys	Phe 235	Glu	Gln	Tyr	Glu	Tyr 240	
	gga Gly															768
	aat Asn															816
_	ccc Pro	-				_	_	-		_			taa			858
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- 400)	=		_			•									
)> 55 Ser		Glu	Lys 5	Leu	Lys	Met	Ser	Phe 10	Glu	Pro	Thr	Val	Ile 15	Glu	
His	Leu	Gly	Val 20	Lys	Met	Tyr	Ser	His 25	Thr	Val	Pro	Ala	Ile 30	Ala	Glu	
Leu	Ile	Ala 35_		Ala	Tyr	Asp	Ala 40	Cys	Ala	Thr	Glu	Val 45	Glu	Val	Arg	
Leu	Phe 50	Asp	Lys	Pro	Glu	His 55	Lys	Ile	Val	Ile	Lys 60	Asp	Asn	Gly	Ile	,
Gly 65	Met	Ser	Phe	Asp	Glu 70	Ile	Asn	Asp	Phe	Tyr 75	Leu	Arg	Ile	Gly	Arg 80	
Asn	Arg	Arg	Glu	Glu 85	Lys	Gln	Ala	Ser	Pro 90	Cys	Gly	Arg	Ile	Pro 95	Thr	
Gly	Lys	Lys	Gly 100	Leu	Gly	Lys	Leu	Ala 105	Leu	Phe	Arg	Leu	Gly 110	Asn	Lys	
Ile	Glu	Ile 115	Ser	Thr	Ile	Gln	Gly 120	Asn	Glu	Arg	Val	Thr 125	Phe	Thr	Leu	
Asp	Tyr 130	Ala	Glu	Ile	Lys	Lys 135	Ser	Glu	Arg	Ile	Tyr 140	Gln	Pro	Glu	Phe	
Gln 145	Lys	Glu	Ser	Val	Lys 150	Pro	Asn	Thr	Glu	Asn 155	Gly	Thr	Thr	Ile	Thr 160	
Leu	Thr	Glu	Leu	Thr 165	Lys	Lys	Gln	Gly	Tyr 170	Pro	Leu	Asp	Asn	Tyr 175	Val	
Gly	His	Leu	Ser 180	Arg	Leu	Phe	Asp	Phe 185	Pro	Ala	Gln	Asp	Phe 190	Lys	Ile	-
Lys	Val	Ser 195	Leu	Asn	Gly	Ser	Glu 200	Pro	Arg	Ile	Ile	Asp 205	Gly	Asn	Leu	
Lys	Tyr 210	Asn	Leu	Val	Thr	Pro 215	Gln	Phe	Glu	Trp	Glu 220	Tyr	Gln	Asp	Leu	
Ala 225	Thr	Asn	Ile	Ser	Ser 230	Leu	Ser	Ser	Lys	Phe 235	Glu	Gln	Tyr	Glu	Tyr 240	

Ser Gly Leu Ile Gln Gly s Phe Ile Thr Thr Glu Lys Pro I Ly 245

Asn Asn Met Lys Gly Ile Thr Leu Phe Ala Asn Gly Arg Met Val Asn 260 265 270

Met Pro Glu Phe Phe Thr Asp Ser Glu Ser Ser His Phe 275 280 285

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gcc agc gaa atc gcc tat cgc ttt gta ttc gga att gaa acc tta ccg $_{20}$ 90 Ala Ser Glu Ile Ala Tyr Arg Phe Val Phe Gly Ile Glu Thr Leu Pro $_{20}$ 25 30

gct gca aaa atg gcg gaa acg ttt gcg ctg aca ttt atg att gct gcg 144
Ala Ala Lys Met Ala Glu Thr Phe Ala Leu Thr Phe Met Ile Ala Ala
35 40 45

ttt ttc gcg ttc agc atg att gcc aac aat gtg cat tac gcg gtt tat
Phe Phe Ala Phe Ser Met Ile Ala Asn Asn Val His Tyr Ala Val Tyr
65 70 75 80

caa agc tgg atg acg ggt att aac tat tgg ctg atg ctg aaa gag gtt 288 Gln Ser Trp Met Thr Gly Ile Asn Tyr Trp Leu Met Leu Lys Glu Val 85 90 95

acc gaa gtc ggc agc gcg gcg tcg atg ttg gat aag ttg tgg ctg 336 Thr Glu Val Gly Ser Ala Gly Ala Ser Met Leu Asp Lys Leu Trp Leu 100 105 110

cct gct ttg tgg ggc gtg gcg gaa gtc atg ttg ttt tgc agc ctt gcc 384
Pro Ala Leu Trp Gly Val Ala Glu Val Met Leu Phe Cys Ser Leu Ala
115 120 125

aag ttc cgc cgt aag acg cat ttt tct gcc gat ata ctg ttt gcc ttc 432 Lys Phe Arg Arg Lys Thr His Phe Ser Ala Asp Ile Leu Phe Ala Phe 130 135 140

cta atg ctg atg att ttc gtg cgt tcg ttc gac acg aaa caa gag cac
Leu Met Leu Met Ile Phe Val Arg Ser Phe Asp Thr Lys Gln Glu His
145 150 155 160

ggt att tcg ccc aaa ccg aca tac agc cgc atc aaa gcc aat tat ttc 528 Gly Ile Ser Pro Lys Pro Thr Tyr Ser Arg Ile Lys Ala Asn Tyr Phe
165 170 175

agc Ser	ttc Phe	ggt Gly	tat Tyr 180	ttt Phę	gtc Val	ggg	cgc Arg	gtg Val 185	ttg Leu	ccg Pro	tat Tyr	cag Gln	ttg Leu 190	F	gat Asp	576
					gtg Val											624
					aat Asn											672
					ttg Leu 230											720
			-	_	caa Gln	_	-		_	_					-	768
					atg Met											816
					aac Asn											864
Asn					gcc Ala											912
					aac Asn 310											960
			-		ctg Leu		_	_	_							1008
	_	Asn	_	Pro	gat Asp	Glu	Lys	Leu	Leu	Pro	Leu	Phe	Asp	Lys		1056
	-	_	_		agg Arg					_						1104
					gca Ala											1152
_	_	_			gat Asp 390	_		_							_	1200
	_				gta Val			-	-		_	_		-		1248
		-		-	tat Tyr			_			_			_		1296
					ggc											1344

440 ctg gtt ttg tac agc ccg t aag gcc gtg caa cag gct gcc a 1392 Leu Val Leu Tyr Ser Pro Asp Lys Ala Val Gln Gln Ala Ala Asn Gln 455 460 gct ttt gcg cct tgc gag att gcc ttc cat cag cag ctt tca acg ttc Ala Phe Ala Pro Cys Glu Ile Ala Phe His Gln Gln Leu Ser Thr Phe 470 475 ctg att cac acg ttg ggc tac gat atg ccg gtt tca ggt tgt cgc gaa 1488 Leu Ile His Thr Leu Gly Tyr Asp Met Pro Val Ser Gly Cys Arg Glu 485 490 ggc tcg gta aca ggc aac ctg att acg ggc gat gca ggc agc ttg aac 1536 Gly Ser Val Thr Gly Asn Leu Ile Thr Gly Asp Ala Gly Ser Leu Asn 500 505 att cgc aac ggc aag gcg gaa tat gtt tat ccg caa taa 1575 Ile Arg Asn Gly Lys Ala Glu Tyr Val Tyr Pro Gln <210> 57 <211> 524 <212> PRT <213> Neisseria gonorrhoeae <400> 57 Met Lys Lys Ser Leu Phe Val Leu Phe Leu Tyr Ser Ser Leu Leu Thr Ala Ser Glu Ile Ala Tyr Arg Phe Val Phe Gly Ile Glu Thr Leu Pro 25 Ala Ala Lys Met Ala Glu Thr Phe Ala Leu Thr Phe Met Ile Ala Ala Leu Tyr Leu Phe Ala Arg Tyr Lys Ala Ser Arg Leu Leu Ile Ala Val Phe Phe Ala Phe Ser Met Ile Ala Asn Asn Val His Tyr Ala Val Tyr Gln Ser Trp Met Thr Gly Ile Asn Tyr Trp Leu Met Leu Lys Glu Val Thr Glu Val Gly Ser Ala Gly Ala Ser Met Leu Asp Lys Leu Trp Leu Pro Ala Leu Trp Gly Val Ala Glu Val Met Leu Phe Cys Ser Leu Ala Lys Phe Arg Arg Lys Thr His Phe Ser Ala Asp Ile Leu Phe Ala Phe 130 135 Leu Met Leu Met Ile Phe Val Arg Ser Phe Asp Thr Lys Gln Glu His 150

Gly Ile Ser Pro Lys Pro Thr Tyr Ser Arg Ile Lys Ala Asn Tyr Phe

Ser Phe Gly Tyr Phe Val Gly Arg Val Leu Pro Tyr Gln Leu Phe Asp 180 185 190

Leu Ser Lys Ile Pro Val Phe Lys Gln Pro Ala Pro Ser Lys Ile Gly

195 200 209

e Val Leu Ile Met Gly Glu Ser d Gln Gly Ser Ile Gln Asi Ala Ala His Leu Lys Leu Phe Gly Tyr Gly Arg Glu Thr Ser Pro Phe 230 Leu Thr Arg Leu Ser Gln Ala Asp Phe Lys Pro Ile Val Lys Gln Ser 250 Tyr Ser Ala Gly Phe Met Thr Ala Val Ser Leu Pro Ser Phe Phe Asn 265 Val Ile Pro His Ala Asn Gly Leu Glu Gln Ile Ser Gly Gly Asp Thr 280 Asn Met Phe Arg Leu Ala Lys Glu Gln Gly Tyr Glu Thr Tyr Phe Tyr Ser Ala Gln Ala Glu Asn Gln Met Ala Ile Leu Asn Leu Ile Gly Lys Lys Trp Ile Asp His Leu Ile Gln Pro Thr Gln Leu Gly Tyr Gly Asn 325 330 Gly Asp Asn Met Pro Asp Glu Lys Leu Leu Pro Leu Phe Asp Lys Ile Asn Leu Gln Gly Arg His Phe Ile Val Leu His Gln Arg Gly Ser 360 His Ala Pro Tyr Gly'Ala Leu Leu Gln Pro Gln Asp Lys Val Phe Gly Glu Ala Asp Ile Val Asp Lys Tyr Asp Asn Thr Ile His Lys Thr Asp 390 Gln Met Ile Gln Thr Val Phe Glu Gln Leu Gln Lys Gln Pro Asp Gly Asn Trp Leu Phe Ala Tyr Thr Ser Asp His Gly Gln Tyr Val Arg Gln Asp Ile Tyr Asn Gln Gly Thr Val Gln Pro Asp Ser Tyr Ile Val Pro Leu Val Leu Tyr Ser Pro Asp Lys Ala Val Gln Gln Ala Ala Asn Gln Ala Phe Ala Pro Cys Glu Ile Ala Phe His Gln Gln Leu Ser Thr Phe 470 Leu Ile His Thr Leu Gly Tyr Asp Met Pro Val Ser Gly Cys Arg Glu 485 Gly Ser Val Thr Gly Asn Leu Ile Thr Gly Asp Ala Gly Ser Leu Asn 505 Ile Arg Asn Gly Lys Ala Glu Tyr Val Tyr Pro Gln

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<212> DNA

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225	230	235	240
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-		gcg gcg ttg ctt ttg Ala Ala Leu Leu Leu 270	
		att tat gcc cac gcc Ile Tyr Ala His Ala 285	
		gcg gca ggt acg ttt Ala Ala Gly Thr Phe 300	
		gcg att tcc aaa gac Ala Ile Ser Lys Asp 315	
		tat ctg cat att gcc Tyr Leu His Ile Ala 335	
	2 3 3 3 3	ttg agt acg gac gct Leu Ser Thr Asp Ala 350	
		att acc tcg cag gcg Ile Thr Ser Gln Ala 365	
		ttg atc ggc agt atc Leu Ile Gly Ser Ile 380	
		ttg tgg atg ggt ttg Leu Trp Met Gly Leu 395	
	Met Gly Lys His Ile	cgt tat tcg ttt ttc Arg Tyr Ser Phe Phe 415	
		agt tcg ata gcg gca Ser Ser Ile Ala Ala 430	
atc gtg cct ctg ccg Ile Val Pro Leu Pro 435			1314
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Leu Leu Thr Glu Lys 20	Val Ser Pro Ile Ile 25	e Ala Leu Ile Leu Val 30	Pro

Leu Ile Gly Ala Leu Leurala Gly Phe Asp Val Ser Gln Leu Phe Tyr Ser Gly Gly Thr Lys Ser Val Thr Gln Ile Val Ile Met Phe Met Phe Ser Ile Leu Phe Phe Gly Ile Met Asn Asp Val Gly Leu Phe Arg Pro Met Ile Gly Gly Leu Ile Lys Leu Thr Arg Gly Asn Ile Val Ala Val Ser Val Gly Thr Val Leu Val Ser Val Val Ala Gln Leu Asp 105 Gly Ala Gly Ala Thr Thr Phe Leu Ser Val Val Pro Ala Leu Leu Pro 120 Leu Tyr Lys Arg Leu His Met Asn Pro Tyr Leu Leu Phe Leu Leu Leu 135 Thr Ser Ser Ala Gly Leu Ile Asn Leu Leu Pro Arg Gly Gly Pro Ile 150 155 Gly Arg Val Ala Ser Val Leu Gly Ala Asp Val Gly Glu Leu Tyr Lys 165 170 Pro Leu Leu Thr Val Gln Ile Ile Gly Val Val Phe Ile Leu Val Leu ____ 180 185 Ser Leu Phe Leu Gly Val Arg Glu Lys Arg Arg Ile Val Arg Glu Leu 200 Gly Ala Leu Pro Ala Val Ala Asp Leu Ile Lys Pro Ala Pro Leu Ser 215 Glu Glu Glu Gln Lys Leu Ala Arg Pro Lys Leu Phe Trp Trp Asn Val 230 Leu Leu Phe Leu Ala Ala Met Ser Leu Leu Phe Ser Gly Ile Phe Pro 250 Pro Gly Tyr Val Phe Met Leu Ala Ala Thr Ala Ala Leu Leu Leu Asn 265 Tyr Arg Ser Pro Gln Glu Gln Met Glu Arg Ile Tyr Ala His Ala Gly 280 Gly Ala Val Met Met Ala Ser Ile Ile Leu Ala Ala Gly Thr Phe Leu 295 Gly Ile Leu Lys Gly Ala Gly Met Leu Asp Ala Ile Ser Lys Asp Leu 310 315 Val His Ile Leu Pro Asp Ala Leu Leu Pro Tyr Leu His Ile Ala Ile Gly Val Leu Gly Ile Pro Leu Glu Leu Val Leu Ser Thr Asp Ala Tyr 345 Tyr Phe Gly Leu Phe Pro Ile Val Glu Gln Ile Thr Ser Gln Ala Gly Val Ala Pro Glu Ala Ala Gly Tyr Ala Met Leu Ile Gly Ser Ile Val Gly Thr Phe Val Thr Project Ser Pro Ala Leu Trp Met Gly I Gly 385 395 400

Leu Ala Lys Leu Ser Met Gly Lys His Ile Arg Tyr Ser Phe Phe Trp 405 410 415

Ala Trp Gly Leu Ser Leu Ala Ile Leu Val Ser Ser Ile Ala Ala Gly 420 425 430

Ile Val Pro Leu Pro 435

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<213> Neisseria gonorrhoeae

<220> <221> CDS <222> (1)..(1152)

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1 5 . 10 15

act tat_ctg tat caa aag ccc aag ctc ttt aaa gga gcg gtt cgg aat 96
Thr Tyr Leu Tyr Gln Lys Pro Lys Leu Phe Lys Gly Ala Val Arg Asn
20 25 30

ctc gaa gcc gca tct tgt aaa tat atc aac gag ata tac caa cga gca 14 Leu Glu Ala Ala Ser Cys Lys Tyr Ile Asn Glu Ile Tyr Gln Arg Ala 35 40 45

gac cca acc gca ccg ctg ttt cat ctg cgt aaa aaa ggc gca atc gtt 192
Asp Pro Thr Ala Pro Leu Phe His Leu Arg Lys Lys Gly Ala Ile Val
50 55 60

cct aaa gaa gaa tac gtc gaa agt ttc gac gat ttg ggc aaa act cgc 240 Pro Lys Glu Glu Tyr Val Glu Ser Phe Asp Asp Leu Gly Lys Thr Arg
65 70 75 80

tac cgt ttt att aaa tcc gtt atc tac gaa cat atg aag aat ggt gcg 288
Tyr Arg Phe Ile Lys Ser Val Ile Tyr Glu His Met Lys Asn Gly Ala
85 90 95

tcg tta gtc tat aac cat att aac aac gag ccg ttt tca gac cat atc 336 Ser Leu Val Tyr Asn His Ile Asn Asn Glu Pro Phe Ser Asp His Ile 100 105 110

gcc cgt caa gtc gcc cgc ttt gcc ggc gca cat act att gtt agt gga 384 Ala Arg Gln Val Ala Arg Phe Ala Gly Ala His Thr Ile Val Ser Gly 115 120 125

tat ctt gct ttt ggc agc gac gaa tct tat aaa aac cat tgg gat acc 432
Tyr Leu Ala Phe Gly Ser Asp Glu Ser Tyr Lys Asn His Trp Asp Thr
130 135 140

cgc gat gtg tat gcc atc cag ctt ttc ggc aag aaa cgt tgg caa ctt 480 Arg Asp Val Tyr Ala Ile Gln Leu Phe Gly Lys Lys Arg Trp Gln Leu 145 150 155 160

act gcc cct gat ttc cct atg cca ttg tat atg caa cag act aaa gat 528

act gat att tcc att cct gaa cat atc gat atg gat att atc ctt gaa . Thr Asp Ile Ser Ile Pro Glu His Ile Asp Met Asp Ile Ile Leu Glu . 180 . 190	576 624												
	624												
gca ggt gat gtc ctc tac atc cca cgc ggt tgg tgg cac aga cct atc Ala Gly Asp Val Leu Tyr Ile Pro Arg Gly Trp Trp His Arg Pro Ile 195 200 205													
ccg ctc ggc tgt gaa acc ttc cac ttc gct gtc ggt acc ttc cca cca Pro Leu Gly Cys Glu Thr Phe His Phe Ala Val Gly Thr Phe Pro Pro 210 215 220	672												
aac ggc tat aat tac ctc gag tgg cta atg aag aaa ttt ccc acc ata Asn Gly Tyr Asn Tyr Leu Glu Trp Leu Met Lys Lys Phe Pro Thr Ile 225 230 235 240	720												
gaa agt ctg cgc cac agt ttc tca gac tgg gag caa gat agg acg cgt Glu Ser Leu Arg His Ser Phe Ser Asp Trp Glu Gln Asp Arg Thr Arg 245 250 255	768												
atc aac gat act gcc gca caa att gct gcc atg att gcc gac ccc gtc Ile Asn Asp Thr Ala Ala Gln Ile Ala Ala Met Ile Ala Asp Pro Val 260 265 270	816												
aat tat gaa gcc ttc agt gaa gac ttt ctc ggc aaa gaa cgt acc gat Asn Tyr Glu Ala Phe Ser Glu Asp Phe Leu Gly Lys Glu Arg Thr Asp 275 280 285	864												
acc gct ttt cat ctc gaa cag ttc gcg aat ccc aac gct act ccg ctt Thr Ala Phe His Leu Glu Gln Phe Ala Asn Pro Asn Ala Thr Pro Leu 290 295 300	912												
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ttg ggg aaa aaa gtg tta gaa cac atc ggt aag aat gaa ccg tta ttg Leu Gly Lys Val Leu Glu His Ile Gly Lys Asn Glu Pro Leu Leu 340 345 350	1056												
ttg aaa aat cta ctg gtt aac ttc aat cag gca aaa cat gaa gaa gtt Leu Lys Asn Leu Leu Val Asn Phe Asn Gln Ala Lys His Glu Glu Val 355 360 365	1104												
agg aag ttg atc tat cag ttg ata gag tta gat ttt ctg gaa att ttg Arg Lys Leu Ile Tyr Gln Leu Ile Glu Leu Asp Phe Leu Glu Ile Leu 370 375 380	1152												
tga 115													
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<400> 61

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Thr Tyr Leu Tyr Gln Lys Pro Lys Leu Phe Lys Gly Ala Val Aman Leu Glu Ala Ala Ser Cys Lys Tyr Ile Asn Glu Ile Tyr Gln Arg Ala Asp Pro Thr Ala Pro Leu Phe His Leu Arg Lys Lys Gly Ala Ile Val 55 Pro Lys Glu Glu Tyr Val Glu Ser Phe Asp Asp Leu Gly Lys Thr Arg Tyr Arg Phe Ile Lys Ser Val Ile Tyr Glu His Met Lys Asn Gly Ala Ser Leu Val Tyr Asn His Ile Asn Asn Glu Pro Phe Ser Asp His Ile 105 Ala Arg Gln Val Ala Arg Phe Ala Gly Ala His Thr Ile Val Ser Gly 120 Tyr Leu Ala Phe Gly Ser Asp Glu Ser Tyr Lys Asn His Trp Asp Thr Arg Asp Val Tyr Ala Ile Gln Leu Phe Gly Lys Lys Arg Trp Gln Leu 150 Thr Ala Pro Asp Phe Pro Met Pro Leu Tyr Met Gln Gln Thr Lys Asp 165 170 Thr Asp Ile Ser Ile Pro Glu His Ile Asp Met Asp Ile Ile Leu Glu 185 Ala Gly Asp Val Leu Tyr Ile Pro Arg Gly Trp Trp His Arg Pro Ile Pro Leu Gly Cys Glu Thr Phe His Phe Ala Val Gly Thr Phe Pro Pro 2,10 Asn Gly Tyr Asn Tyr Leu Glu Trp Leu Met Lys Lys Phe Pro Thr Ile Glu Ser Leu Arg His Ser Phe Ser Asp Trp Glu Gln Asp Arg Thr Arg Ile Asn Asp Thr Ala Ala Gln Ile Ala Ala Met Ile Ala Asp Pro Val Asn Tyr Glu Ala Phe Ser Glu Asp Phe Leu Gly Lys Glu Arg Thr Asp Thr Ala Phe His Leu Glu Gln Phe Ala Asn Pro Asn Ala Thr Pro Leu Ser Asp Asp Val Arg Leu Arg Leu Asn Ala Asn Asn Leu Asp Thr Leu Glu Lys Gly Tyr Leu Ile Gly Asn Gly Met Lys Ile Ser Val Asp Glu Leu Gly Lys Lys Val Leu Glu His Ile Gly Lys Asn Glu Pro Leu Leu Leu Lys Asn Leu Leu Val Asn Phe Asn Gln Ala Lys His Glu Glu Val

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cgc caa acc agc ctg acg ggt aaa gtg att ctg aca cga ccg ttg tca Arg Gln Thr Ser Leu Thr Gly Lys Val Ile Leu Thr Arg Pro Leu Ser 20 25 30	96													
ttt tcc cta tgg acg aca ttt gca tcg ata tct gcg tta ttg att atc Phe Ser Leu Trp Thr Thr Phe Ala Ser Ile Ser Ala Leu Leu Ile Ile 35 40 45	144													
ctg ttt ttg ata ttt ggt aac tat acg cga aag aca aca gtg gag gga Leu Phe Leu Ile Phe Gly Asn Tyr Thr Arg Lys Thr Thr Val Glu Gly 50 55 60	192													
caa att tta cct gca tcg ggc gta atc agg gtg tat gca ccg gat acg Gln Ile Leu Pro Ala Ser Gly Val Ile Arg Val Tyr Ala Pro Asp Thr 65 . 70 . 75 . 80	240													
ggg aca att aca gcg aaa ttc gtg gaa gat gga gaa aag gtt aag gct Gly Thr Ile Thr Ala Lys Phe Val Glu Asp Gly Glu Lys Val Lys Ala 85 90 95	288													
ggc gac aag cta ttt gcg ctt tcg acc tca cgt ttc ggc gca gga gat Gly Asp Lys Leu Phe Ala Leu Ser Thr Ser Arg Phe Gly Ala Gly Asp 100 105 110	336													
agc gtg cag cag ttg aaa acg gag gca gtt ttg aag aaa acg ttg Ser Val Gln Gln Leu Lys Thr Glu Ala Val Leu Lys Lys Thr Leu 115 120 125	384													
gca gaa cag gaa ctg ggt cgt ctg aag ctg ata cac ggg aat gaa acg Ala Glu Gln Glu Leu Gly Arg Leu Lys Leu Ile His Gly Asn Glu Thr 130 135 140	432													
cgc agc ctt aaa gca act gtc gaa cgt ttg gaa aac cag gaa ctc cat Arg Ser Leu Lys Ala Thr Val Glu Arg Leu Glu Asn Gln Glu Leu His 145 150 155 160	480													
att tcg caa cag ata gac ggt cag aaa agg cgc att aga ctt gcg gaa Ile Ser Gln Gln Ile Asp Gly Gln Lys Arg Arg Ile Arg Leu Ala Glu 165 170 175	528													
gaa atg ttg cag aaa tat cgt ttc cta tcc gcc aat gat gca gtg cca Glu Met Leu Gln Lys Tyr Arg Phe Leu Ser Ala Asn Asp Ala Val Pro 180 185 190 ·	576													
aaa caa gaa atg atg aat gtc aag gca gag ctt tta gag cag aaa gcc Lys Gln Glu Met Met Asn Val Lys Ala Glu Leu Leu Glu Gln Lys Ala	624													

195 200 205

aaa ctt gat gcc tac' cgd a gaa gaa gtc ggg ctg ctt cag gatc 67 Lys Leu Asp Ala Tyr Arg Arg Glu Glu Val Gly Leu Leu Gln Glu Ile 210 215 220

cgc acg cag aat ctg aca ttg gcc agc ctc ccc caa gcg gca tga 71
Arg Thr Gln Asn Leu Thr Leu Ala Ser Leu Pro Gln Ala Ala
225 230 235

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<211> 236 <212> PRT

<213> Neisseria meningitidis

<400> 63

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Arg Gln Thr Ser Leu Thr Gly Lys Val Ile Leu Thr Arg Pro Leu Ser 20 25 30

Phe Ser Leu Trp Thr Thr Phe Ala Ser Ile Ser Ala Leu Leu Ile Ile 35 40 45

Leu Phe Leu Ile Phe Gly Asn Tyr Thr Arg Lys Thr Thr Val Glu Gly 50 55 60

Gln Ile Leu Pro Ala Ser Gly Val Ile Arg Val Tyr Ala Pro Asp Thr
65 - 70 75 80

Gly Thr Ile Thr Ala Lys Phe Val Glu Asp Gly Glu Lys Val Lys Ala . 90 95

Gly Asp Lys Leu Phe Ala Leu Ser Thr Ser Arg Phe Gly Ala Gly Asp 100 105 110

Ser Val Gln Gln Leu Lys Thr Glu Ala Val Leu Lys Lys Thr Leu 115 120 125

Ala Glu Glu Leu Gly Arg Leu Lys Leu Ile His Gly Asn Glu Thr 130 140

Arg Ser Leu Lys Ala Thr Val Glu Arg Leu Glu Asn Gln Glu Leu His 145 150 155 160

Ile Ser Gln Gln Ile Asp Gly Gln Lys Arg Arg Ile Arg Leu Ala Glu 165 170 175

Glu Met Leu Gln Lys Tyr Arg Phe Leu Ser Ala Asn Asp Ala Val Pro $180\ \cdot \ 185$

Lys Gln Glu Met Met Asn Val Lys Ala Glu Leu Leu Glu Gln Lys Ala 195 200 205

Lys Leu Asp Ala Tyr Arg Glu Glu Val Gly Leu Leu Gln Glu Ile 210 215 220

Arg Thr Gln Asn Leu Thr Leu Ala Ser Leu Pro Gln Ala Ala 225 230 235

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<210> 65 <211> 229 <212> PRT <213> Neisseria gonorrhoeae

Ala Tyr Gly Arg Glu Glu Ala Gly Leu Leu Gln Glu Ile Arg Thr Gln
20 25 30

Asn Leu Thr Leu Ala Ser Leu Pro Lys Arg His Glu Thr Glu Gln Ser 35 40 45

Gln Leu Glu Arg Thr Met Ala Asp Ile Ser Gln Glu Val Leu Asp Phe 50 55 60

Glu Met Arg Ser Glu Gln Ile Ile Arg Ala Gly Arg Ser Gly Tyr Ile 65 70 75 80

Ala Ile Pro Asn Val Glu Val Gly Arg Gln Val Asp Pro Ser Lys Leu 85 90 95

Leu Leu Ser Ile Val Pro Glu Arg Thr Glu Leu Tyr Ala His Leu Tyr 100 105 110

Ile Pro Ser Ser Ala Ala Gly Phe Ile Lys Pro Lys Asp Lys Val Val 115 120 125

Leu Arg Tyr Gln Ala Tyr Pro Tyr Gln Lys Phe Gly Leu Ala Ser Gly 130 135 140

Ser Val Val Ser Val Ala Lys Thr Ala Leu Gly Arg Gln Glu Leu Ser 145 150 155 160

Gly Leu Gly Met Val Ser Ser Asp Leu Ala Lys Ser Asn Glu Pro Val 165 170 175

Tyr Leu Val Lys Ile Lys Pro Asp Lys Pro Thr Ile Thr Ala Tyr Gly
180 185 190

Glu Glu Lys Pro Leu Gln Ile Gly Met Thr Leu Glu Ala Asp Ile Leu 195 200 205

His Glu Lys Arg Arg Leu Tyr Glu Trp Val Leu Glu Pro Ile Tyr Ser 210 215 220

Met Ser Gly Arg Leu 225

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<211> 924

<212> DNA

<213> Neisseria gonorrhoeae

<220>

<221> CDS

<222> (1)..(921)

<400> 66

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			ggg Gly 20													96
			gat Asp													144
			tcc Ser													192
			ctc Leu													240
	-	-	cgc Arg	_				_	_							288
		-	agc Ser 100		-		-	-			-			-		336
			tct Ser					_	_	_	_			_		384
			ctc Leu	_	_	-	_	_		_	-					432
	_	_	att Ile			_				-			-			480
			gac Asp													528
		-	ccg Pro 180		_							_		_		576
_			ttg Leu					_		-			-		_	624
			gaa Glu													672
			tac Tyr													720
			acc Thr													768
_			atc Ile		_		_	_	_		-		-		_	816

gat gtt gta gaa gtt gc c tct tac gac caa tcc gac att 864 Asp Val Val Glu Val Ala Pro Ser Tyr Asp Gln Ser Asp Ile Thr Ala 280 ttg gcc ggc gcc aca att gcc ttg gaa atg ctt tac ctt caa ggt gcg 912 Leu Ala Gly Ala Thr Ile Ala Leu Glu Met Leu Tyr Leu Gln Gly Ala 295 300 924 aaa aag gac tga Lys Lys Asp 305 <210> 67 <211> 307 <212> PRT <213> Neisseria gonorrhoeae <400> 67 Met Gln Tyr Ser Thr Leu Ala Gly Gln Thr Asp Asn Ser Leu Val Ser 10 Asn Asn Phe Gly Phe Leu Arg Leu Pro Leu Asn Phe Met Pro Tyr Glu Ser His Ala Asp Trp Val Ile Thr Gly Val Pro Tyr Asp Met Ala Val 40 Ser Gly Arg Ser Gly Ala Arg Phe Gly Pro Glu Ala Ile Arg Arg Ala Ser Val Asn Leu Ala Trp Glu His Arg Arg Phe Pro Trp Thr Phe Asp Val Arg Glu Arg Leu Asn Ile Ile Asp Cys Gly Asp Leu Val Phe Ser Phe Gly Asp Ser Arg Asp Phe Val Glu Lys Met Glu Ala His Ala Gly 105 Lys Leu Leu Ser Phe Gly Lys Arg Cys Leu Ser Leu Gly Gly Asp His 120 Phe Ile Thr Leu Pro Leu Leu Arg Ala His Ala Arg Tyr Phe Gly Lys 135 Leu Ala Leu Ile His Phe Asp Ala His Thr Asp Thr Tyr Asp Asn Gly 155 150 Ser Glu Tyr Asp His Gly Thr Met Phe Tyr Thr Ala Pro Lys Glu Gly Leu Ile Asp Pro Ser Arg Ser Val Gln Ile Gly Ile Arg Thr Glu His 185 Ser Lys Lys Leu Pro Phe Thr Val Leu Ser Ala Pro Lys Val Asn Glu 200 Asp Ser Val Glu Glu Thr Val Arg Lys Ile Lys Glu Thr Val Gly Asn 215 Met Pro Val Tyr Leu Thr Phe Asp Ile Asp Cys Leu Asp Pro Ser Phe

Ala Pro Gly	Thr Gly 245	Thr Pro	Val Cy	s Gly 250	Gly	Leu	Ser	Ser	Asp	Arg	
Ala Leu Lys	Ile Leu 260	Arg Gly	Leu Th 26		Leu	Asp	Ile	Val 270	Gly	Met	
Asp Val Val 275	Glu Val	Ala Pro	Ser Ty 280	r Asp	Gln	Ser	Asp 285	Ile	Thr	Ala	
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Lys Lys Asp 305											
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cca cta tcc Pro Leu Ser 50											192
tgg cgg gcg Trp Arg Ala 65	Ala Gln										240
cag cgt gat Gln Arg Asp											288
ctt ccc cat Leu Pro His				r Tyr							336
att tct tcc Ile Ser Ser 115											384
acc tta ttt Thr Leu Phe 130		-	-			_		_			432
gat acg cag Asp Thr Gln 145											480

				gcc Ala 165											gac Asp	528
				cat His												576
	-		-	gct Ala						-	-			_	_	624
				aaa Lys												672
				gag Glu												720
				agc Ser 245												768
	_	_		ctg Leu		_	_	_	_		_	_	-	-		816
				tta Leu												864
_	_		_	agc Ser		_				-	-	_		-		912
				tct Ser												960
				aat Asn 325												1008
				ttg Leu												1056
ggc Gly	aaa Lys	atc Ile 355	cat His	gaa Glu	gcc Ala	gaa Glu	gcg Ala 360	caa Gln	tac Tyr	ggg	gcc Ala	gcc Ala 365	gaa Glu	gca Ala	cag Gln	1104
				gag Glu												1152
				gcg Ala												1200
				cgt Arg 405												1248
tac	ggc	atc	cgc	aac	cgg	ctg	gaa	gta	ata	cgg	gcg	cgg	cag	gaa	gtc	1296

Tyr Gly Ile Arg Asn A	Arg Leu Glu Val Ile 425	e Arg Ala Arg Gln 430	Glu Val
gcc caa gca gaa cag a Ala Gln Ala Glu Gln L 435			
gct tat ttg cgc ttg g Ala Tyr Leu Arg Leu V 450			
ttt gcg gaa taa Phe Ala Glu 465			1404
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Cys Leu Thr Leu Thr P	Pro Tyr Leu Gln His 25	Glu Leu Phe Ser 3	Ala Met
Lys Ser Tyr Phe Ser L	Lys Tyr Ile Leu Pro 40	o Val Ser Leu Phe ' 45	Thr Leu
Pro Leu Ser Leu Ser P 50	Pro Ser Val Ser Ala 55	Phe Thr Leu Pro 6	Glu Ala
Trp Arg Ala Ala Gln G 65	Gln His Ser Ala Asp 70	Phe Gln Ala Ser : 75	His Tyr 80
Gln Arg Asp Ala Val A	Arg Ala Arg Gln Glr 90	-	Ala Phe 95
Leu Pro His Val Ser A 100	Ala Asn Ala Ser Tyr 105	Gln Arg Gln Pro 110	Pro Ser
Ile Ser Ser Thr Arg G	Glu Thr Gln Gly Trg 120	Ser Val Gln Val	Gly Gln
Thr Leu Phe Asp Ala A	Ala Lys Phe Ala Glr 135	n Tyr Arg Gln Ser . 140	Arg Phe
Asp Thr Gln Ala Ala G 145	Glu Gln Arg Phe Asp 150	o Ala Ala Arg Glu 155	Glu Leu 160
Leu Leu Lys Val Ala G 165	Glu Ser Tyr Phe Asr 170		Arg Asp 175
Thr Val Ala Ala His A 180	Ala Ala Glu Lys Glu 185	Ala Tyr Ala Gln 190	Gln Val
Arg Gln Ala Gln Ala I 195	Leu Phe Asn Lys Gly 200	y Ala Ala Thr Ala 205	Leu Asp
Ile His Glu Ala Lys A 210	Ala Gly Tyr Asp Ası 215	n Ala Leu Ala Gln 220	Glu Ile
Ala Val Leu Ala Glu I 225 2	Lys Gln Thr Tyr Glu 230	ı Asn Gln Leu Asn 235	Asp Tyr 240

Thr	Asp	Leu	Asp	Ser 245		n	Ile	Glu	Ala 250	Ile	Asp	Thr	Ala		Leu	
Leu	Ala	Arg	Tyr 260	Leu	Pro	Lys	Leu	Glu 265	Arg	Tyr	Ser	Leu	Asp 270	Glu	Trp	
Gln	Arg	Ile 275	Ala	Leu	Ser	Asn	Asn 280	His	Glu	Tyr	Arg	Met 285	Gln	Gln	Leu	
Ala	Leu 290	Gln	Ser	Ser	Gly	Gln 295	Ala	Leu	Arg	Ala	Ala 300	Gln	Asn	Ser	Arg	
Tyr 305	Pro	Thr	Val	Ser	Ala 310	His	Val	Gly	Tyr	Gln 315	Asn	Asn	Leu	Tyr	Thr 320	
Ser	Ser	Ala	Gln	Asn 325	Asn	Asp	Tyr	His	Tyr 330	Arg	Gly	Lys	Gly	Met 335	Ser	
Val	Gly	Val	Gln 340	Leu	Asn	Leu	Pro	Leu 345	Tyr	Thr	Gly	Gly	Glu 350	Leu	Ser	
Gly	Lys	Ile 355	His	Glu	Ala	Glu	Ala 360	Gln	Tyr	Gly	Ala	Ala 365	Glu	Ala	Gln	
Leu	Thr 370	Ala	Thr	Glu	Arg	His 375	Ile	Lys	Leu	Ala	Val 380	Arg	Gln	Ala	Tyr	
Thr 385		Ser	Gly	Ala	Ala 390	Arg	Tyr	Gln	Ile	Met 395	Ala	Gln	Glu	Arg	Val 400	
Leu	Glu	Ser	Ser	Arg 405	Leu	Lys	Leu	Lys	Ser 410	Thr	Glu	Thr	Gly	Gln 415	Gln	
Tyr	Gly	Ile	Arg 420	Asn	Arg	Leu	Glu	Val 425	Ile	Arg	Ala	Arg	Gln 430	Glu	Val	٠
Ala	Gln	Ala 435	Glu	Gln	Lys	Leu	Ala 440	Gln	Ala	Arg	Tyr	Lys 445	Phe	Met	Leu	
Ala	Tyr 450	Leu	Arg	Leu	Val	Lys 455	Glu	Ser	Gly	Leu	Gly 460	Leu	Glu	Thr	Val	
Phe 465	Ala	Glu														
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	_			-				-		_		ggc Gly		_		96

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gtg ggc gct ttg Val Gly Ala Leu 50		l Val Phe		_	-	
ttc ccg cgc gtt Phe Pro Arg Val 65						
gcc cta tat ttg Ala Leu Tyr Leu				-		-
ata gtc ggt tcg Ile Val Gly Ser 100		_	_			
gtc ggc aat ctt Val Gly Asn Leu 115		-				
att ttt ggc ttg Ile Phe Gly Leu 130		Lys Tyr				
gct gac gta aaa Ala Asp Val Lys 145 -			-		-	
tta ttg act ttg Leu Leu Thr Leu						
gat aaa gat tgg Asp Lys Asp Trp 180		-		-		
gac ctg tat tac Asp Leu Tyr Tyr 195						
gcg cgc cca cat Ala Arg Pro His 210		r Ser Lys	_	-		
ccg cca act tgc Pro Pro Thr Cys 225	-					696
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Thr Leu Gly Ile 20	Cys Ala Le	u Leu Ala 25	Phe Cys	Phe Gly	Ala Ala 30	Ile
Ala Ser Gly Tyr	His Leu Gl	u Tyr Glu	Tyr Gly	Tyr Arg	Tyr Ser	Ala

l Val Phe Leu Leu Leu Leu Ala . Val Gly Ala Leu Ala Se Phe Pro Arg Val Ser Ser Val Val Leu Leu Ile Tyr Val Gly Thr Thr 70 Ala Leu Tyr Leu Pro Val Gly Trp Leu Tyr Gly Ala Pro Ser Tyr Gln 90 Ile Val Gly Ser Ile Leu Glu Ser Asn Pro Ala Glu Ala Arg Glu Phe 105 Val Gly Asn Leu Pro Gly Ser Leu Tyr Phe Val Gln Ala Leu Phe Phe 120 Ile Phe Gly Leu Thr Val Trp Lys Tyr Cys Val Ser Val Gly Val Phe 135 Ala Asp Val Lys Asn Tyr Lys Arg Arg Ser Lys Ile Trp Leu Thr Ile 150 Leu Leu Thr Leu Ile Leu Ser Cys Ala Val Met Glu Lys Ile Ala Gly 170 165 Asp Lys Asp Trp Arg Glu Pro Asp Ala Gly Leu Leu Asn Ile Phe 185 Asp Leu Tyr Tyr Asp Leu Ala Phe Arg Ala Gly Thr Ile Cys Arg Gln 195 200 Ala Arg Pro His Phe Gly Ser Ser Lys Ser Val Asn Met Ala Tyr 215 220 Pro Pro Thr Cys Ala Gln Val 225 <210> 72 <211> 2607 <212> DNA <213> Neisseria meningitidis <220> <221> CDS <222> (1)..(2604) <400> 72 atg gct gcc aac caa cgt tac cgc aaa ccg ctg ccc ggt acg gat ttg Met Ala Ala Asn Gln Arg Tyr Arg Lys Pro Leu Pro Gly Thr Asp Leu 1 10 gaa tac tac gac gcg cgt gcg gcg tgt gag ggc atc aaa ccc ggc tct Glu Tyr Tyr Asp Ala Arg Ala Ala Cys Glu Gly Ile Lys Pro Gly Ser 20 tac gac aag ctg cct tac acg agc cgc att ttg gcg gag aat ttg gtc Tyr Asp Lys Leu Pro Tyr Thr Ser Arg Ile Leu Ala Glu Asn Leu Val 35 40 aac cgc gcg gac aaa gtc gat ttg ccg acg ctg caa agc tgg ctg ggt Asn Arg Ala Asp Lys Val Asp Leu Pro Thr Leu Gln Ser Trp Leu Gly 50 55

cag c Gln I 65						cag l.n								ccd	gcg Ala 80	240
cgg g Arg V																288
gca g Ala G																336
gtg a Val A																384
gtg g Val G	-	-				-		_			_					432
atc of Ile 0 145	•	-	_	_		-	_	_								480
aaa a Lys T		-														528
atg o Met H																576
aac g Asn G			-			-		_	-		_	_	_		_	624
cca c Pro H																672
gaa g Glu A 225		_		_		_		_			_		_	_		720
gat a Asp I																768
gcg a Ala T																816
gtg g Val V	_				_	-				_			_	_	_	864
tct a Ser]			_	_						_	-	_				912
gcg a Ala 1 305		_		_		-		-					_		_	960
aaa d Lys I																1008

				325					330					335	,	
gcc Ala								gat Asp 345							tat Tyr	1056
ccg Pro																1104
ggc Gly	_	_		-			-				_	-	_	_		1152
aaa Lys 385																1200
gac Asp		-				-					-					1248
aat Asn	_	_		-	-	-	-		-							1296
cgc Arg																1344
ggt Gly																1392
gaa Glu 465																1440
tgt Cys																1488
atc Ile														cgc Arg		1536
ttt Phe	_		-			_				_	-			gct Ala		1584
														ttc Phe		1632
														cgc Arg		1680
														gcc Ala 575		1728
														ttc Phe		1776

				caa Gln											cca Pro	1824
				atc Ile												1872
				tta Leu												1920
				gac Asp 645												1968
_	_	-		gaa Glu		_	-		-		_		-	_	-	2016
				gca Ala			_		-		_		-		-	2064
				aat Asn												2112
				cgc Arg												2160
		-	_	atg Met 725		-	-		_			_		_		2208
				atc Ile												2256
				gca Ala												2304
-	-	-		ttc Phe		_			_			_			-	2352
				ctg Leu												2400
	_	_		acg Thr 805	-			-	-	_		_	_			2448
_	_	_	-	acc Thr		_			_						_	2496
_	-			acc Thr	_	_		_		-	_	-		_	_	2544
	-	-		ggc Gly	-	_				-	_	-		_	_	2592

865

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<211> 868

<212> PRT

<213> Neisseria meningitidis

<400> 73

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1 10 15

Glu Tyr Tyr Asp Ala Arg Ala Ala Cys Glu Gly Ile Lys Pro Gly Ser 20 25 30

Tyr Asp Lys Leu Pro Tyr Thr Ser Arg Ile Leu Ala Glu Asn Leu Val 35 40 45

Asn Arg Ala Asp Lys Val Asp Leu Pro Thr Leu Gln Ser Trp Leu Gly 50 60

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Lys Thr Ala Phe Glu Asn Val Asp Val Ile Pro Ala Gly Asn Gly Ile 165 170 175

Met His Gln Ile Asn Leu Glu Lys Met Ser Pro Val Val Gln Val Lys 180 185 190

Asn Gly Val Ala Phe Pro Asp Thr Cys Val Gly Thr Asp Ser His Thr 195 200 205

Pro His Val Asp Ala Leu Gly Val Ile Ser Val Gly Val Gly Leu 210 215 220

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				_	-	-		ggc Gly	_				-			384
_		_				_		aac Asn							-	432
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ttg Leu 225																720
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